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Result
No.
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Maximum DB seq
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Database
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Maximum Match 100%
Listing first 45 s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score
    protein search, using sw
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        length: 0
length: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Issued Patents AA:*

1: /EMC Celerra_SIDS3/ptodata/2/iaa/5_COMB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/2/iaa/6_COMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/iaa/7_COMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/iaa/H_COMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/iaa/PCTUS_COMB.pep:*

6: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*

7: /EMC_Celerra_SIDS3/ptodata/2/iaa/backfiles1.pep:*
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97
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
    ВB
                   US-09-920-262A-6
US-09-674-973A-64
US-09-674-973A-64
US-09-260-582-47
US-08-260-582-47
US-08-260-582-47
US-08-260-582-47
US-08-260-5471-48
US-09-685-548A-126
US-09-685-548A-126
US-09-685-474-79-22
US-08-661-479-22
US-08-661-479-22
US-08-661-479-22
US-08-661-479-22
US-08-661-479-20
US-09-641-803-18
US-09-641-801-18
US-09-641-801-19
US-08-482-018-10
US-08-482-018-10
US-08-482-018-10
US-08-482-018-10
US-09-188-001-10
US-09-188-001-10
US-09-188-001-10
US-09-188-001-10
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81.494 Million cell updates/sec
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6, Appli
14, Appl
47, Appl
47, Appl
48, Appl
126, Appl
22, Appl
22, Appl
22, Appl
366, Appl
314, Appl
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316, Appl
317, Appl
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| 45                   | 44                   | 43                  | 42                  | 41              | 40               | 39               | 38                 | 37                 | 36                 | 35                 | 34                 | υ<br>U             | 32                | 31               | 30               | 29               | 28               | 27               |
|----------------------|----------------------|---------------------|---------------------|-----------------|------------------|------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|-------------------|------------------|------------------|------------------|------------------|------------------|
| 26                   | 26                   | 26                  | 26                  | 26              | 27               | 27               | 27                 | 27                 | 27                 | 27                 | 27                 | 27                 | 27                | 28               | 28               | 28               | 28               | 28               |
| 26.8                 |                      | 26.8                |                     |                 |                  |                  |                    |                    |                    | •                  |                    | ٠                  | ٠                 | ٠                |                  | 28.9             | •                | 28.9             |
| 8                    | œ                    | œ                   | œ                   | υt              | 18               | 18               | 18                 | 18                 | 18                 | 18                 | 18                 | 18                 | 11                | 18               | 18               | 18               | 16               | <b>T</b> 5       |
| w                    | ω                    | w                   | w                   | N               | ۲                | ۲                | ۲                  | ۲                  | -                  | _                  | ۲                  | Н                  | N                 | ۲                | μ                | ۳                | N                | ۲                |
| US-09-641-528B-23996 | US-09-641-528B-16358 | US-09-641-528B-8847 | US-09-641-528B-1718 | US-09-788-006-9 | US-09-017-205-49 | US-09-017-205-48 | US-08-421-695A-101 | US-08-421-698A-100 | US-08-421-697A-100 | US-08-421-696A-100 | US-08-303-052A-100 | US-08-421-702A-100 | US-09-809-517A-21 | US-08-443-977-30 | US-08-443-976-30 | US-08-084-718-30 | US-10-044-708A-3 | US-08-080-0/3-26 |
| Sequence             | Sequence             | Sequence            | Sequence            | Sequence        | Sequence         | Sequence         | Sequence           | Sequence           | Sequence           | Seguence           | Sequence           | Sequence           | Seguence          | Sequence         | Sequence         | Sequence         | Sequence         | sequence         |
| 23996, A             | 16358, A             | 8847, Ap            | 1718, Ap            | 9, Appli        | 49, Appl         | •                |                    | 100, App           |                    | 100, App           |                    |                    |                   | •                | -                | •                | ъ                | בקלה יסי         |

# ALIGNMENTS

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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Shealy, David

APPLICANT: Knight, David

APPLICANT: Kcaight, David

APPLICANT: Scallon, Bernie

APPLICANT: Giles-Komar, Jill

APPLICANT: Peritt, David

TITLE OF INVENTION: ANTI-IL-12 ANTIBODIES, COMPOSITIONS, MET

FILE REFERENCE: CEN0248

CURRENT APPLICATION NUMBER: US/09/920,262A

CURRENT FILING DATE: 2000-08-07

PRIOR APPLICATION NUMBER: 60/236,358

PRIOR APPLICATION NUMBER: 60/236,827

PRIOR FILING DATE: 2000-09-29

NUMBER: OF SEQ ID NOS: 15

NUMBER: OF SEQ ID NOS: 15
Sequence 64, Application US/09674973A
Patent No. 6759046
GENERAL INFORMATION:
APPLICANT: NO. 67590468k Hydro ASA
TITLE OF INVENTION: Peptides
FILE REFERENCE: 26625-296
CURRENT APPLICATION UMWBER: US/09/674,973A
CURRENT FILING DATE: 2001-06-04
NUMBER OF SEQ ID NOS: 459
SOFTWARE: PatentIn version 3.0
SEQ ID NO 64
LENGTH: 15
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US-09-920-262A-6
                                                                                                                                                                                                                                                   US-09-674-973A-64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: Homo sapiens US-09-920-262A-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver 3.1
SEQ ID NO 6
LENGTH: 9
TYPE: PRT
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Best Local
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83.3%;
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Pred. No. 5e+05;
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; SEQ ID NO 14
; LENGTH: 11
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-307-265A-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Gu, Trent
APPLICANT: Orita, Satoshi
APPLICANT: Orita, Satoshi
APPLICANT: Han, Min
TITLE OF INVENTION: RAS SUPPRESSOR SUR-5
FILE REFERENCE: UTC-03732
CURRENT APPLICATION NUMBER: US/09/307,265A
CURRENT FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 16
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US-09-307-265A-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity bz...
Thes 5; Conservative
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/260,582
PILING DATE: 16-JUN-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 31,544
REFERENCE/DOCKET NUMBER: GI 5236
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 14, Application US/09307265A
Patent No. 6225456
GENERAL INFORMATION:
APPLICANT: Gu, Trent
APPLICANT: Orita, Satoshi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 47, Application US/08260582 Patent No. 5635182
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                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: MCCOY, John M.
APPLICANT: Lu, Zhijian
TITLE OF INVENTION: METHOD OF DETECTING LIGAND
TITLE OF INVENTION: INTERACTIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   y Match 33.0%;
Local Similarity 54.5%;
hes 6; Conservative
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                                                                                                                                                                                                                                                                                          COUNTRY:
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62.5%;
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Pred. No. 1e+02;
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; Patent No. 5635182;
; GENERAL INFORMATION:
; APPLICANT: McCoy, John M.
; APPLICANT: Lu, Zhijian
; TITLE OF INVENTION: METHOD OF DETECTIONS
; TITLE OF INVENTION: INTERACTIONS
; NUMBER OF SEQUENCES: 76
; CORRESPONDENCE ADDRESS:
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                                                                                                US-08-260-582-48
                                                                                                                                                                                           TELEFAX: (61/)
INFORMATION FOR SEQ ID NO: 48
SEQUENCE CHARACTERISTICS:
FRIGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-260-582-48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ANTI-SENSE: NO US-08-260-582-47
                                                                                                                                                                                                                                                                      COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/260,582
FILING DATE: 16-JUN-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 31,544
REFERENCE/DOCKET NUMBER: G1 5236
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                               Best Local Similarity
Matches 5; Conserv
                                                               Query Match
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Best Local Similarity 62.5%;
Matches 5; Conservative
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TELEFAX: (617) 876-58
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 02140
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: TRM PC compati
                                                                                                               MOLECULE TYPE: protein HYPOTHETICAL: NO ANTI-SENSE: NO
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LENGTH: 12 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
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5635182
11 PYTAPGKO 18
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87 CambridgePark Drive
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                                 Conservative
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                                                32.0%;
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                                                                                                                                                                                                                                             48:
                                 2;
                                                Score 31; DB 1;
Pred. No. 1.2e+02;
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Pred. No. 1.2e+02;
                                 Mismatches
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                                                               Length 12;
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PWLAPGEQ 11

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; MOLECULE TYPE:
; HYPOTHETICAL: N
; ANTI-SENSE: NO
PCT-US95-05471-48
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RESULT 8
US-09-865-548A-126
; Sequence 126, Application US/09865548A
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                                                                                                                                                  Query Match 32.0%;
Best Local Similarity 62.5%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 48, Application PC/TUS9505471 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/05471
INFORMATION FOR SEQ ID NO: 47:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:

TITLE OF INVENTION: METHOD OF DETECTING LIGAND INTERA
NUMBER OF SEQUENCES: 76

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: METHOD OF DETECTING LIGAND INTERACTIONS NUMBER OF SEQUENCES: 76
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
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Local Similarity 62.5%;
es 5; Conservative
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                                                                                    PWLAPGEO 11
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                                                                                                                                                    ; Score 31; DB
; Pred. No. 1.2e
2; Mismatches
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Pred. No. 1.2e+02;
2; Mismatches 1
                                                                                                                                                                      DB 5;
1.2e+02;
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                                                                                                                                                                                      Length 12;
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Applicar, Ricardo
APPLICANT: Choe, Sunghwa
APPLICANT: Feldmann, Kenneth A.
TITLE OF INVENTION: DWF4 POLYNUCLEOTIDES, POLYPEPTIDES AND USES THEREOF
FILE REFERENCE: 11996-070001
CURRENT APPLICATION NUMBER: US/09/502,426B
CURRENT FILING DATE: 2000-02-11
PRIOR APPLICATION NUMBER: US 60/119,657
PRIOR FILING DATE: 1999-02-11
PRIOR APPLICATION NUMBER: US 60/119,658
PRIOR FILING DATE: 1999-02-11
PRIOR APPLICATION NUMBER: US 60/119,658
PRIOR FILING DATE: 1999-02-11
NUMBER OF SEQ ID NOS: 30
SOPTWARR: FastSEQ for Windows Version 4.0
SEQ ID NO 27
LENGTH: 14
RESULT 10
US-08-333-565-22
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                                                                                                                                       Query Match
Best Local Similarity
Thehes 5; Conservi
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                                                                                                                                                                                                                                   ; OTHER INFORMATION: Heme binding domain US-09-502-426B-27
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US-09-502-426B-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Admon, Arie
TITLE OF INVENTION: METHOD OF IDENTIFYING PEPTIDES CAPABLE OF BINDING TO MHC MOLECULE:
TITLE OF INVENTION: PEPTIDES IDENTIFIED THEREBY AND THEIR USES
FILE REPERENCE: 01/22080
CURRENT APPLICATION NUMBER: US/09/865,548A
CURRENT FILING DATE: 2001-05-16
PRIOR APPLICATION NUMBER: US 60/290,958
PRIOR FILING DATE: 2001-05-16
NUMBER OF SEQ ID NOS: 204
SOFTWARE: PACENTIN VERSION 3.1
SEQ ID NO 126
FENCETU: 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 27, Application US/09502426B Patent No. 6987025
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Best Local (
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APPLICANT: Beer, Ilan
APPLICANT: Ziv, Tamar
                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
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55.6%;
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Pred. No. 2.1e+02
                                                                                                                                                           Mismatches
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Sequence 22, Application US/08333565 Patent No. 5622852 GENERAL INFORMATION: APPLICANT: KORSMEYER, Stanley J.

KORSMEYER, Stanley J.

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US-08-661-479-22
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INFORMATION FOR SEQ ID NO: 22
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
STRANDENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22, Application US/08661479 Patent No. 5834209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION: APPLICANT: KORSME
              APPLICATION NUMBER: US/08/661,479
FILING DATE: 11-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/333,565
FILING DATE: 31-OCT-1994
                                                                                                                                                                                                            ZIP: 94301
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Bcl-x/Bcl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: 1i
ATTORNEY/AGENT INFORMATION:
                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15726A-000700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                              STREET: 379 Lytton Avenue CITY: Palo Alto STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/333,565
FILING DATE: 31-OCT-1994
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
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ENTION: Bcl-x/Bcl-2 ASSOCIATED CELL DEATH
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linear
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                                                                                                    US/08/661,479
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Pred. No. 2.5e+02;
0; Mismatches 1
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US-09-042-353-366
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Matches 5; Conserv
                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/810,279
FILING DATE: 17-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/853,408
FILING DATE: 18-MAR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-JUN-1992
PRIOR APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
PRIOR APPLICATION NUMBER: US 08/053,131
APPLICATION NUMBER: US 08/053,131
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/053,131
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APPLICATION NUMBER: US 08/053,131
PRIOR APPLICATION DATA:
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APPLICANT: Lonberg, Nils
APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 6255458-Human Animals for
TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 421
CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP
ADDRESSE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (415) 326-24 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/042,353 FILING DATE: 13-MAR-1998 CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15726A-000700
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                           APPLICATION NUMBER: US 08/096,762
FILING DATE: 22-JUL-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/155,301
FILING DATE: 18-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
STRANDEDNESS: single
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TELEPHONE: (415)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 366, Application US/09042353
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Two Embarcadero Center, Eighth Floor
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326-2422
NO: 22:
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US 08/161,739
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Pred. No. 2.5e+02;
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RESULT 13
US-08-758-417A-214
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STRANDEDNESS:
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-09-042-353-366
                                                                                                                                                                                                                                                                                                                                                                   Sequence 214, Application US/08758417A Patent No. 6300129
GENERAL INFORMATION:
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Best Local
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INFORMATION FOR SEQ ID NO: 366:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PILING DATE: 10-OCT-1996
PRIOR APPLICATION DATA:

APPLICATION UMMBER: US 08/758
FILING DATE: 02-DEC-1996
PRIOR APPLICATION UMMBER: WO PCT/US
APPLICATION UMMBER: WO PCT/US
FILING DATE: 01-DEC-1997
ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 0146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 576-0200
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/728,463

FILING DATE: 10-OCT-1996

PRIOR APPLICATION NUMBER: WO PCT/US96/16433

FILING DATE: 10-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08/352,322 FILING DATE: 07-DEC-1994 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/165,699 FILING DATE: 10-DEC-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
          COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                      APPLICANT: Lonberg, Nils
Kay, Robert M.
Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 6300129-Human Animals for
Producing Heterologous Antibodies
                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and
STREET: Two Embarcadero
  CURRENT
                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 l Similarity
5; Conserva
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                                                                                                                                                               CITY: San Francisco
STATE: California
                                                                                                                        ZIP: 94111-3834
                                                                                                                                            COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 29; DB 2; Length 9; Pred. No. 5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   014643-009040US
                                                                                                                                                                                                      Townsend and Crew LLP
Center, Eighth Floor
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US-08-278-774-20
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                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20, Application US/08278774 Patent No. 6653450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Berg, Richard A
APPLICANT: Toman, David P
APPLICANT: Wallace, Donald
TITLE OF INVENTION: MUTATED RECOMBINANT COLLAGENS
NUMBER OF SEQUENCES: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (415) 576-03
INFORMATION FOR SEQ ID NO: 214:
SEQUENCE CHARACTERISTICS:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                             ZIP: 94303
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                 STREET: 2500 Fal
CITY: Palo Alto
STATE: Californ
                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 214:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Serafini, Andrew T.
REGISTRATION NUMBER: 41,303
REFERENCE/DOCKET NUMBER: 01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: <U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08/728,463
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                                                                                                                                                                   California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 03-DEC-1993
APPLICATION NUMBER: US 08/155,301
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 08/209,741 FILING DATE: 09-MAR-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 18-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/096,762
                                                                                                                                                                                                        2500 Faber Place
                                                                                                                                                USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                             COLLAGEN CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29.9%; Score 29; DB 2; 83.3%; Pred. No. 5e+05;
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                       Version #1.25
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Length 9; 1; Indels

0

Gaps

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Sequence 1, Application Us/09856920

Patent No. 6740325

GENERAL INFORMATION:

APPLICANT: Yedd Research and Development Co.

PITLE OF INVENTION: Peptide-based vaccine for influenza

FILE REFERENCE: 9822 PCT

CURRENT APPLICATION NUMBER: Us/09/856,920

CURRENT FILING DATE: 2002-02-19

NUMBER OF SEQ ID NOS: 9

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 1

LENGTH: 18

Type: PAT

ORGANISM: Influenza virus

US-09-856-920-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX:

INFORMATION FOR SEQ ID NO: 20
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-278-774-20
Search completed: May 30, 2006, 15:09:26 Job time : 20.3333 secs
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US-09-856-920-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/278,774
FILING DATE: 22-JUL-1994
CLASSIFICATION: 435
ATTORNEY, AGENT INFORMATION:
NAME: Rafayko, Kathi L
REGISTRATION NUMBER: 36,644
REFERENCE/DOCKET NUMBER: 94-018
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 354-4642
TELEPAX: (415) 354-4752
                                                                             5 SNCYPYDVP 13
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Maximum Match
Listing first
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Sequence:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq
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                                                                                                                                                                                                                                 length:
length:
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                                                                                                                                                                                                                                                                                                                    283416 seqs, 96216763 residues
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96
                                                                  PIR 80:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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Biocceleration Ltd.
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SUMMARIES

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R;Kimura, A.; Takata, M.; Fukushi, Y.; Mori, H.; Matsui, H.; Chiba, S. Biosci. Biotchem. 61, 1091-1098, 1997
A;Title: A catalytic amino acid and primary structure of active site in Aspergillus nige A;Reference number: PT0090; MUID:97399878; PMID:9255970
A;Accession: PT0090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Fonagy, A.; Schoofs, L.; Proost, P.; Van Damme, J.; De Loof, A. Comp. Biochem. Physiol. C 103, 135-142, 1992
A;Title: Isolation and primary structure of two sulfakinin-like peptides from the fleshf A;Reference number: A56632; MUID:93083101; PMID:1360367
A;Accession: A56632
                                                                                                      A;Cross-references: UNIPROT:Q7M3M5; UNIPARC:UPI000017BF08C;Keywords: glycosidase; hydrolase
                                                                                                                                                                                                                                                                                                                                                  alpha-glucosidase (EC 3.2.1.20) - honeybee (fragment)
C;Species: Apis mellifera (honeybee)
C;Date: 07-Jull1997 #sequence_revision 29-Aug-1997 #text_change 09-Jul-2004
C;Accession: PT0090
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                                                                                                                                                                  A; Molecule type: protein A; Residues: 1-15 < KIM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 F;14/Modified site: amidated carboxyl end (Phe)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C; Keywords: amidated carboxyl end; neuropeptide; sulfoprotein F;1-14/Product: neosulfakinin-I #status experimental <NSKl>F;6-14/Product: neosulfakinin-I #status experimental <NSKl>F;9/Binding site: sulfate (Tyr) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Note: sequence extracted from NCBI backbone (NCBIP: 120391)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:P41493; UNIPARC:UPI000013054D A;Experimental source: heads
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: protein A; Residues: 1-14 < FON>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Accession: A56632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N;Contains: neosulfakinin-I (Neb-SK-I)
C;Species: Sarcophaga bullata
C;Date: 21-Jul-1995 #sequence_revision 21-Jul-1995 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N;Alternate names: Neb-SK-II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     neosulfakinin-II - flesh fly (Sarcophaga bullata)
                            Query Match
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h 25.0%;
Similarity 57.1%;
4; Conservative
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Pred. No. 6.1e+02;
2; Mismatches 2;
  Score 24; DB 2;
Pred. No. 9.9e+02;
2; Mismatches 1
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N;Alternate names: LSK
C:Species: Leucophaea maderae (Madeira cockroach)
C;Date: 17-Mar-1987 #sequence_revision 17-Mar-1987 #text_change 09-Jul-2004
C;Accession: A01622
R;Nachman, R.J.; Holman, G.M.; Haddon, W.F.; Ling, N.
Science 234, 71-73, 1986
A;Title: Leucosulfakinin, a sulfated insect neuropeptide with homology to gastr.
A;Reference number: A01622; MUID:86315858; PMID:3749893
A;Accession: A01622
A;Molecule type: protein
A;Residues: 1-1 cANC>
A;Cross-references: UNIPROT:P04428; UNIPARC:UPI000012E960
C;Superfamily: gastrin
C;Keywords: amidated carboxyl end; hormone; sulfoprotein
F;6/Binding site: sulfate (Tyr) (covalent) #status experimental
F;11/Modified site: amidated carboxyl end (Phe) #status experimental
perisulfakinin - American cockroach
C;Species: Periplaneta americana (American cockroach)
C;Decies: Action 14-May-1993 #sequence_revision 14-May-1993 #text_change 09-Jul-2004
C;Accession: A60656
R;Veenstra, J.A.
Neuropeptides 14, 145-149, 1989
A;Title: Isolation and structure of two gastrin/CCK-like neuropeptides from A;Reference number: A60656; MUID:90137190; PMID:2615921
A;Accession: A60656
A;Molecule type: protein
A;Residues: 1-11 <VEE>
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C;Keywords: extracellular protein; hydrolase; serine prot
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                                                             structure of two gastrin/CCK-like neuropeptides from the Americar 0656; MUID:90137190; PMID:2615921
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57.1%;
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Pred. No. le+03;
1; Mismatches
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Pred. No. 1.2e+03;
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gasawara, N.; Yasunaga, T.; Kuhara
DNA Res. 8, 11-22, 2001
A;Title: Complete genome sequence
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A;Title: Dominant T-cell-receptor beta chain variable region V beta A;Reference number: A49655; MUID:94068553; PMID:8248215
A;Accession: D49655
                                                                                                                                                                                        A;Cross-references: UNIPROT:Q8X8T5; UNIPARC:UPI00000D08D2; GB:BA000007; PIDN:BAB36242.1; A;Experimental source: strain O157:H7, substrain RIMD 0509952
                                                                                                                                                                                                                                                                                                                                                                                                                    his operon leader peptide [imported] - Eschetichia coli (strain 0157:H7, substrain C;Species: Escherichia coli C;Species: 18-Jul-2001 #sequence_revision 18-Jul-2001 #text_change 09-Jul-2004
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A;Residues: 1-16 <GRO>
A;Residues: 1-16 <GRO>
A;Cross-references: DIPARC:UPI000017C3CA
A;Experimental source: peripheral blood lymphocytes
A;Note: sequence extracted from NCBI backbone (NCBIP:140448)
C;Keywords: T-cell receptor
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D49655
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C;Comment: This neuropeptide stimulates hindgut contractions.
C;Keywords: amidated carboxyl end; neuropeptide; sulfoprotein
F;6/Binding site: sulfate (Tyr) (Covalent) #status experimental
F;11/Modified site: amidated carboxyl end (Phe) #status experimental
                                                                                                                                                      A;Gene:
                                                                                                                                                                                                                                                A; Status: preliminary A; Molecule type: DNA
                                                                                                                                                                                                                                                                                      A;Reference number: A99629;
A;Accession: C90981
                                                                                                                                                                                                                                                                                                                                                              R;Hayashi, T.; Makino, K.; Ohnishi, M.; Kurokawa, F
gasawara, N.; Yasunaga, T.; Kuhara, S.; Shiba, T.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T-cell-receptor beta chain variable region, TCR V beta (clone 2) - human (f. C;Species: Homo sapiens (man)
C;Date: 07-Apr-1994 #sequence_revision 18-Nov-1994 #text_change 30-May-1997
C;Accession: D49655
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                                                                        Score 23; DB 2;
Pred. No. 1.6e+03;
0; Mismatches 1
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Pred. No. 1.
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Shinagawa,
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napin large chain L1A - Swedish turnip (fragments)
C;Species: Brassica napus var. rapifera (Swedish turnip, rutabaga)
C;Date: 19-Mar-1998 #sequence_revision 17-Apr-1998 #text_change 31-Dec-2004
C;Accession: $70340
R;Neumann, G.M.; Condron, R.; Thomas, I.; Polya, G.M.
Biochim. Biophys. Acta 1295, 34-43, 1996
A;Title: Purification and sequencing of multiple forms of Brassica napus seed napin larg
A;Reference number: $70340; MUID:96283791; PMID:8679671
A;Accession: $70340
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                                                                                                                                                                                                                                                                                                                                                 R;Mooser, G.; Hefta, S.A.; Paxton, R.J.; Shively, J.E.; Lee, T.D. J. Biol. Chem. 266, 8916-8922, 1991
A;Title: Isolation and sequence of an active-site peptide contain A;Reference number: A39841; MUID:91224988; PMID:1827439
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A;Residues: 1-12;13-18 <NEU>
A;Cross-references: UNIPROT:039344; UNIPROT:Q9S9F0; UNIPARC:UPI000017B02E; UNIPARC:UPI00
C;Superfamily: Alpha amylase inhibitor
                                                                                                                                                                                           A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-9 <MOO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Date: 20-Mar-1992 #sequence_revision 20-Mar-1992 #text_change 09-Jul-2004 C;Accession: B39841
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dextransucrase (EC 2.4.1.5) - Streptococcus sobrinus (fragment) C; Species: Streptococcus sobrinus
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A; Residues: 1-16 <STO>
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                                                                                                                ;Cross-references: UNIPROT:Q7M0L7; UNIPARC:UPI000017AC69;Keywords: glycosyltransferase; hexosyltransferase
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36.4%;
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    22.9%;
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Pred. No. 1.8e+03;
3; Mismatches 4
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Pred. No. 1.6e+03;
0; Mismatches 1
    Score 22;
Pred. No.
    DB 2; Length 9;
2.8e+05;
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RESULT 13
A34818
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C;Species: Ovis orientalis ariee, Ovis ammon aries (domesti
C;Date: 19-Mar-1997 #sequence_revision 01-Aug-1997 #text_ch
C;Accession: S29379
R;Reiersen, H.; Sletten, K.; McKinley-McKee, J.S.
Eur. J. Biochem. 211, 861-869, 1993
A;Title: Affinity labelling of sorbitol dehydrogenase from A;Reference number: S29379; MUID:93170323; PMID:8436142
A;Accession: S29379
                                          C;Species: Cajanus cajan (pigeon pea)
C;Date: 13-Jul-1990 #sequence_revision 13-Jul-1990 #text_change
C;Accesion: A34818
R;Mawal, Y.R.; Mawal, M.R.; Ranjekar, P.K.
Biochem. Biophys. Res. Commun. 166, 1446-1452, 1990
A;Title: Unusual denaturation properties of vicilin from Cajanus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Simmer, J.P.; Kelly, R.E.; Scully, J.L.; Grayson, D.R.; Rinker Jr., A.G.; Bergh, S.T.; Proc. Natl. Acad. Sci. U.S.A. 86, 4382-4386, 1989
A;Title: Mammalian aspartate transcarbamylase (ATCase): sequence of the ATCase domain and A;Reference number: A33917; MUID:89282776; PMID:2543974
A;Accession: A33917
A;Reference number: A34818; MUID:90165956; PMID:2306256
A;Accession: A34818
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C;Species: Cricetulus griseus (Chinese hamster)
C;Date: 09-Mar_1990 #sequence_revision 09-Mar-1990 #text_change 07-Nov-1997
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                                                                                                                                                                                 vicilin 72K chain - pigeon pea (fragment)
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C;Superfamily: alcohol dehydrogenase; long-chain alcohol
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A; Residues: 1-11 <SIM>
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Pred. No. 2.7e+03;
1; Mismatches 3
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Pred. No. 1.5e+03;
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Query Match
Best Local Similarity 3
Warches 3; Conservat!
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C;Species: Aplysia californica (California sea hare)
C;Species: Aplysia californica (California sea hare)
C;Date: 02-Feb-1990 #sequence_revision 02-Feb-1990 #text_change 09-Jul-2004
C;Accession: A30330
R;Lloyd, P.E.; Connolly, C.M.
J. Neurosci. 9, 312-317, 1989
A;Title: Sequence of pedal peptide: a novel neuropeptide from the central nervous system A;Reference number: A30330; MUID:89110403; PMID:2913209
A;Status: preliminary
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PC7075
PC7075
PC7075

Guanylate cyclase (EC 4.6.1.2), soluble, alpha-1 chain - mouse (fragment)
C.Species: Mus musculus (house mouse)
C.Pate: 18-Aug-2000 #sequence_revision 18-Aug-2000 #text_change 09-Jul-2004
C.Accession: PC7075
C.Accession: PC7075
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A;Residues: 1-14 <TSU's
A;Residues: 1-14 <TSU's
A;Cross-references: UNIPROT:Q7M058; UNIPARC:UPI000017C675
A;Experimental source: strain C57BL/6Cr Slc, male; brain, striatum
C;Keywords: brain; phosphorus-oxygen lyase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-reterences: Jan...
C;Keywords: neuropeptide
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A;Molecule type: protein
A;Residues: 1-15 <LLO>
A;Cross-references: UNIPROT:Q7M3P8; UNIPARC:UPI000017BD9A
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A;Title: Proteome analysis of mouse brain: Two-dimensional electrophoresis profiles of tA;Reference number: PC7072
A;Accession: PC7075
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A;Residues: 1-7 <MAW>
A;Cross-references: UNIPARC:UPI000017B040
Search completed: May 30, 2006, 15:08:20 Job time : 13 secs
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

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RESULT 2
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| 8                    | 31.2<br>62.5<br>rative            | ; EAA2074;<br>Hypothet:<br>1778 MW; |                | sequence<br>DDBJ who<br>ata.   | d compa<br>n yoeli<br>(2002).  | er J.C.                                | igh J.,<br>I.R. II  |                                | റ്റെട്  | (LARC                          | lii.<br>Api                         | version 10   | cated i          | PRELIMINARY      |            | 16<br>17<br>17                | 166                | 15                         | 14<br>14                                  | 100        | .i       |
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|                      | +03                               | C_DNA.<br>E109 CR                   | http           | erived<br>n (WGS)  | . et   | h H.<br>ffma                           | oaibi A<br>Sinden 1 | ay M.F.                        | 1038<br>ij T<br>leng                              | NA].                           | porida                              |  | EMBL.            | AA.              |            |                               |                    |                            |   |            |          |
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RESULT 3

QUJUUS_MOUSE

QUJUUS_MOUSE

QUJUUS;

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QUJUUS;

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O1-OCT-2000, integrated into UniProtKB/TrEMBL.

DT

O1-FCB-2006, entry version 1.

DT

O7-FCB-2006 entry version 9.

B-Raf protein (Fragment).

Mus musculus (Mouse).

CL Eukaryota; Metazoa; Chordata; Craniata; Vertebrat Mus musculus (Mouse).

CL Eukaryota; Metazoa; Chordata; Craniata; Vertebrat Musmalia; Eutheria; Euarchontoglires; Glires; Rod Mus musculus (Muridae; Murinae; Mus.

OC Muroidea; Muridae; Murinae; Mus.

OX NCBI_TAXID-10090;

RP NUCLEOTIDE SEQUENCE.

RA Barnier J. V., Papin C., Eychene A., Lecoq O.;

RA Barnier J. V., Papin C., Eychene Multiple protein is specific expression.";

J. Biochem. 270:23381-23389(1995).

RA Barnier J. V.;

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Mus musculus (Mouse).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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                                                                     Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AF025882; AAD22136.1; -; Genomic DNA. EMBL; AF025881; AAD22136.1; JOINED; Genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "A novel human myo-inositol monophosphatase gene, susceptibility region for bipolar disorder."; Mol. Psychiatry 2:393-397(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-2000, sequence version 1.
07-FEB-2006, entry version 9.
Inosine monophosphatase 2 (Fragment).
                                                                                                                                                  Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.
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lgaru M., Karkera J.D., Sharma M., Berrettin
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RESULT 4

QROFT3 C

AC QROFT3 C

AC QROFT3;

DT 01-JUN-2

DT 07-FEB-2

DT 07-FEB-2

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CG Archosau

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QBQFT3
                                                                                             01-NOV-1995, integrated into UniProtKB/Swiss-Prot.
01-NOV-1995, sequence version 1.
07-MAR-2006, entry version 32.
07-MAR-2006, entry version 32.
Neosulfakinin-2 (Neosulfakinin-II) (Neb-SK-II).
Sarcophaga bullata (Grey flesh fly) (Neobellieria bullata).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Oestroidea;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Oestroidea;
Sarcophagidae; Sarcophaga; Neobellieria.
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SEQUENCE
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GO; GO:000584; C:nucleus; RCA.
GO; GO:0005886; C:plasma membrane; IDA.
GO; GO:0005886; C:plasma membrane; IDA.
GO; GO:0003677; F:DNA binding; RCA.
GO; GO:0008713; F:protein-tyrosine kinase activity; RCA.
GO; GO:0008714; F:zinc ion binding; RCA.
GO; GO:000875; P:cellular physiological process; RCA.
GO; GO:0006355; P:regulation of transcription, DNA-deper
NON_TER
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11
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                                                                                                                                                                                                                                                                                                                                                                               P41493;
                                                                                                                                                                                                                                                                                                                                                                                                                      NSK2
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PROTEIN SEQUENCE
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Fonagy A., Schoofs L., Proost P., van Damme J., de Loof A.; "Isolation and primary structure of two sulfakinin-like peptid the fleshfly, Neobellieria bullata."; Comp. Biochem. Physiol. 103C:135-142(1992).

-!- FUNCTION: Myotropic peptide.
-!- SUBCELIULAR LOCATION: Secreted protein.

peptides

MEDLINE=93083101; PubMed=1360367;

rissuE=Head;

PIR; A56632; A56632.

InterPro; IPR013152; Gastrin CCK. InterPro; IPR013259; Sulfakinin.

PEPTIDE Amidation;

PROSITE; PS00259; GASTRIN;

Direct protein sequencing;

Neuropeptide;

Sulfation.

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PF08257; Sulfakinin;

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http://www.uniprot.org/terms

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RESULT
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P83012;
29-MAR-2005,
                      urrect protein sequencing; Heme; Inner membrane; Iron; Metal-binding; Mitochondrion; Oxidoreductase CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-21534268; PubMed=11676495; DOI=10.1006/bbrc.2001.5826; Schuurmans Stekhoven F. M.A.H., Filk G., Wendelaar Bonga S. B.; "N-terminal sequences of small ion channels in rectal glands of sharks: a blochemical hallmark for classification and phylogeny?"; Biochem. Blophys. Res. Commun. 288:670-675 (2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Scyliorhinus canicula (Spotted dogfish) (Spotted catshark). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes; Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Scyliorhinidae;
NCBI_TaxID=7830;
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07-FEB-2006, entry version 15.
Cytochrome c oxidase polypeptide
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MEDLINE=21534268; Pu
                                                                                                               GO; GO:0005739; C:mitochondrion; InterPro; IPR003204; Cyt_c_ox5A.
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                                                                                                                                                                                                                                                                    SIMILARITY: Belongs to the cytochrome c oxidase Va
                                                                                                                                                                                                                                                                                              SUBCELLULAR LOCATION: Mitochondrion; mitochondrial
                                                                                                                                                                                                                                                                                                                                         CATALYTIC ACTIVITY: 4 ferrocytochrome c + O(2)
                                                                                                                                                                                                                                                                                                                                                                  oxidase, the terminal
                                                                                                                                                                                                                                                                                                                                                                                         FUNCTION:
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                                                                                                                                                                                                                                                                                                                                 Blophys. Res. Commun. 288:670-6/5\20041.

TION: This is the heme A-containing chain of cytochrome critical transports of the terminal oxidase in mitochondrial electron transports. A ferrocytochrome c + O(2) = 4 ferricytoch
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                                                                                                                                                                                                                       the UniProt Consortium,
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Pred. No. 9.2e
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neosulfakinin-2.
/FTId=PRO_000043897.
Sulfotyrosine (Potential).
Phenylalanine amide (Potential).
Cytochrome c oxidase polypeptide Va /FTId=PRO_0000195215.
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                                                                                                                                                IDA.
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                                                                                                                                                                                                                                                                  inner membrane. family.
                                                                         Membrane;
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RESULT 8
Q7M3M5 APIME
ID Q7M3M5;
AC Q7M3M5;
DT 15-DEC-2003, s
DT 15-DEC-2006, e
DE Alpha-glucosid
OS Apis mellifera
OC Eukaryota; Met
OC Neoptera; Endo
OC Neoptera; Endo
OC NcBI_TaxID=746
RN [1]
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O12692

D1 O12692

D2 O12692

D1 O12692

D1 O1-JUL-1

DT O1-JUL-1

DT O1-JUL-1

DT O7-MAR-2

DB Game-gag

OS Simian-H

OC Viruses

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RN [1]

RP NUCLEOTI

RC STRAIN=S

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Best Local
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Best Local
   Eukaryota; Metazoa; Arthro
Neoptera; Endopterygota; i
Apidae; Apis.
NCBI_TaxID=7460;
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01-JUL-1997,
07-MAR-2006,
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NON TER
SEQUENCE 18
                                                                                                                                                                     07-FEB-2006, entry version 6.
Alpha-glucosidase (EC 3.2.1.20)
Apis mellifera (Honeybee).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; U89249; AAC57878.1; -; Genomic_DNA. GO; GO:0019013; C:viral nucleocapsid; IEA. GO; GO:0005198; F:strutral molecule activity; Thispers. Temponomy: Te
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "A cell-free stock of simian-human immunodeficiency virus that causes AIDS in pig-tailed macaques has a limited number of amino acid substitutions in both SIVmac and HIV-1 regions of the genome and has
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=97312452; PubMed=9168893; DOI=10.1006/viro.1997.8534;
Stephens E.B., Mukherjee S., Sahni M., Zhuge W., Raghavan R.,
Singh D.K., Leung K.K., Atkinson R.B., Li Z., Joag S.V., Liu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Simian-Human immunodeficiency virus.
Viruses; Retro-transcribing viruses;
Lentivirus; Primate lentivirus group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gag polyprotein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pfam; PF00540; Gag_p17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               offered cytotropism.";
Virology 231:313-321(1997).
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NCBI_TaxID=57667;
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                                                                                                                                                                                                                                              sequence version entry version 6.
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1720
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50.0%;
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57.1%;
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                                                                                                            Hymenoptera;
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Pred. No. 9.8e+03;
2; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 25;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                                                                                           (Fragment).
                                                                                                                                            Hexapoda; Insecta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                            Apocrita; Aculeata;
                                                                                                                                                                                                                                                                                                                                                                                15
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Attribution-NoDerivs License
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RESULT

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RESULT 10
Q4YLT9 PLABE
ID Q4YLT9 P
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DT 05-JUL-2
DT 05-JUL-2
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Q9R563_ECOLI
Q9R563;
                                Q4YLT9;
Q5-JUL-2005, integrated into UniProtKB/TrEMBL.
05-JUL-2005, sequence version 1.
07-FEB-2006, entry version 4.
107-FEB-2006, entry version 4.
Hypothetical protein (Fragment).
ORFNames-PB400723.00.0;
Plasmodium berghei.
Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
NCBI_TaxID=5821;
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MEDLINE=93328296; PubMed=8392972;
Dytoc M., Soni R., Cockerill F.III.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriaceae; Escherichia.
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01-MAY-2000, sequence version 1.
07-FEB-2006, entry version 14.
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"Multiple determinants of verotoxin-producing Escherichia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=562;
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M., Fukushi Y.,
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66.7%;
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Pred. No. 1.4e+04;
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ID QSFBRS S
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DT 15-MAR-2
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CC Mammalia
OC Eukaryot
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OX NCBI_Tax
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15-MAR-2005,
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STRAIN=TKU, WZ, and BAN; TISSUE=Kidney;
PubMed=15897627; DOI=10.1538/expanim.54.173;
Adjei S., Sato A., Tanaka S., Kobayashi E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Name=TXN;
Name=TXN;
Suncus murinus (House shrew) (Musk shrew).
Suncus murinus (House shrew) (Musk shrew).
Translata; Vertebrata; Euteleostomi;
Translata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A., Berriman M., Florens L., Janssen C.S., Pain A., Christophides James K., Rutherford K., Harris B., Harris D., Churcher C.M., Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J., Bidwell S.L., Rajandream M.A., Carucci D.J., Yates J.R. III, Kafacos F.C., Janse C.J., Barrell B.G., Turner C.M.R., Waters
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Crania
Mammalia; Eutheria; Laurasiatheria;
Crocidurinae; Suncus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9378;
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Q5FBR5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transcriptomic, and proteomic analyses."; Science 307:82-86(2005).
                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                        Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                     Exp. Anim. 54:173-180(2005).
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PubMed=15637271; DOI=10.1126/science.1103717;
                                                                                                                                                                                                                                                                                                                                                                                                                          Development and characterization
                                                                                                                                                                                                                                                                                                                                                                                                                                                        shikawa A.;
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                                                                                                                                                                                     AB167756;
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Pred. No. 1.7e+04;
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Science 234:71-73(1986).

-!- FUNCTION: Changes the frequency and amplitude of contractions the hingut. Inhibits muscle contraction of hindgut.
                                                                                                                                                                                                                                               PROTEIN SEQUENCE.

MEDLINE=86315858; PubMed=3749893;

Machman R.J., Holman G.M., Haddon W.F., Ling N.;

"Leucosulfakinin, a sulfated insect neuropeptide with homology"
                                                                                                                                                                                                                                                                                                                                                           Leucophaea maderae (Madeira cockroach).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea, Dictyptera; Blattaria; Blaberoidea;
Blaberidae; Oxyhaloinae; Leucophaea.
                              InterPro; IPR013152; Gastrin CCK.
InterPro; IPR013259; Sulfakinin.
Pfam; PF08557; Sulfakinin; 1.
PROSITE; PS00259; GASTRIN; 1.
                                                                                                                                                                                                                                              gastrin and cholecystokinin.";
                                                                                                PIR; A01622; GMROL
                                                                                                                                 Distributed under the Creative Commons
                                                                                                                                           Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
                                                                                                                                                                          the hingut. Inhibits muscle contraction of hindgut.
-!- SIMILARITY: Belongs to the gastrin/cholecystokinin family.
                 Amidation;
                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=6988;
 Direct protein sequencing; Hormone;

1 11 Leucosulfakinin-1.
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                  Sulfation
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Query Match
Best Local :
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LSCA1 LEUMA STANDARD; PRT;
P04428;
13-AUG-1987, integrated into UniProtKE
13-AUG-1987, sequence version 1.
07-MAR-2006, entry version 36.
Leucosulfakinin-1 (Leucosulfakinin-1)
                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=92155439; PubMed=1786859; DOI=10.1016/0020-711X(91)90133-8; Sinha U., Wolz S.A., Lad P.J.; The sextracellular serine proteases from Streptomyces fradiae."; Int. J. Biochem. 23:979-984(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q7M198_STRFR
Q7M198;
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Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
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07-FEB-2006, entry version 7.
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27.3%;
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Pred. No. 1.7e+04;
4; Mismatches 4
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OCCUPATION 
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ID P78359_HUMAN
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea; Dictyoptera; Blattaria; Blattoidea;
Blattidae; Blattinae; Periplaneta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Isolation and structure of two gastrin/CCK-like neuropeptides from the American cockroach homologous to the leucosulfakinins."; Neuropeptides 14:145-149(1989)
-!- FUNCTION: Stimulates hindgut contractions.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Corpora cardiaca;
MEDLINE=90137190; PubMed=2615921; DOI=10.1016/0143-4179(89)90038-3;
                                                                                  Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR013152; Gastrin_CCK.
InterPro; IPR013259; Sulfakinin.
Pfam; PF08257; Sulfakinin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Distributed under the Creative Commons
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1 11 Perisulfakinin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Title:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OM protein -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9
                                                                                                                                                                                                                                                       DB seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein search, using sw
                                                                                                                                                                                                                                                     g length: 0
g length: 18
                                                                                                           A_Geneseq_8:*
1: geneseqp198
2: geneseqp200
3: geneseqp200
                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-758-165A-11
96
                                                                                                                                                                                                                                                                                                                                    2589679 seqs, 457216429 residues
                                                                                                                                                                                                                                                                                                                                                                    Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                      BLOSUM62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              May 30, 2006, 14:59:31; Search time 61.6667 Seconds
                                                                                                                                                                                          Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                IDGQKVDEQFPQHGLVKQ 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GenCore version 5.1.8
Copyright (c) 1993 - 2006 Biocceleration Ltd.
             geneseqp2003as:*
geneseqp2003bs:*
geneseqp2004s:*
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geneseqp2006s:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (without alignments)
133.458 Million cell updates/sec
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score and is d. No. is the number of results predicted by chance to have a re greater than or equal to the score of the result being printed, is derived by analysis of the total score distribution.

# SUMMARIES

| 23                 | 22       | 21       | 20       | 19       | 18       | 17       | 16                 | 15       | 14                 | 13       | 12       | 11       | 10                 | 9                 | 8                  | 7                  | σ                  | ъ                  | 4                  | w        | 2                  | Ļ                  | Result<br>No. |
|--------------------|----------|----------|----------|----------|----------|----------|--------------------|----------|--------------------|----------|----------|----------|--------------------|-------------------|--------------------|--------------------|--------------------|--------------------|--------------------|----------|--------------------|--------------------|---------------|
| 33                 | 33       | 33       | 33       | 33       | 33       | 33       | 33                 | 33       | 34                 | 34       | 34       | 34       | 34                 | 34                | 35                 | 35                 | 35                 | 36                 | 36                 | 36       | 78                 | 96                 | Score         |
| 34.4               | 34.4     | 34.4     |          | 34.4     |          | 34.4     | 34.4               | 34.4     |                    |          | 35.4     |          | 5                  |                   | ა                  | 36.5               | 6                  | 7                  | 7.                 | 37.5     | ۲                  | 100.0              | Query         |
| 15                 | 15       | 15       | 15       | 15       | 15       | 15       | 13                 | 10       | 18                 | 17       | 15       | 15       | 15                 | 15                | 18                 | 16                 | 16                 | 14                 | 14                 | 14       | 15                 | 18                 | Length        |
| a                  | 6        | 9        | σ        | 6        | σ        | თ        | 9                  | æ        | œ                  | œ        | თ        | σ        | δ                  | N                 | œ                  | 9                  | 9                  | 9                  | 7                  | ر.       | 7                  | 8                  | 80            |
| ABP95372           | ABP94780 | ABP94539 | ABP95117 | ABP94098 | ABP95554 | ABP94247 | AED95486           | ADL98126 | ADR10609           | ADM96833 | ABP95194 | ABP95588 | ABP94693           | AAR50232          | ADR10610           | AEE34193           | AEE33637           | AED79318           | ADG97265           | ABP46438 | ADC64569           | ADR10611           | ID            |
| Abp95372 HLA prote |          | HLA      | HLA      | HLA      | HLA      | HLA      | Aed95486 T cell de | Candida  | Adr10609 Dog IgE e | Tiss     | HLA      | HL.A     | Abp94693 HLA prote | Aar50232 Sequence | Adr10610 Cat IgE e | Aee34193 Wheat gli | Aee33637 Wheat gli | Aed79318 Human B L | Adg97265 scFV VHCD | Humar    | Adc64569 Horse imm | Adr10611 Horse IgE | Description   |

| 00000000000000000000000000000000000000  | 22<br>25<br>27                               |
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|   | 4444   |
| 15<br>15<br>16<br>16<br>11<br>14<br>14<br>15<br>15<br>15  | 15<br>15<br>15                               |
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| ABC70956 ABC70908 ABC70908 ABC34208 ABC834208 ABC834215 AAW15125 AAW98200 ABP46692 ABP97662 ADG97262 ADG97519 ABD79572 ABD79572 ABP94687 ABP94687 ABP94689 ABP94689 ABP95184  | ABR32079<br>ABR32127<br>ADV31901<br>ADV31853 |
| Aec70956 Aec70958 Aec34208 Aec33641 Aew15125 Aaw15125 Aag98200 Abp46692 Abp46692 Adg97262 Adg97519 Aed79315 Aed79315 Aed79315 Aed794689 Abp94689 Abp94689 Abp94689 Abp95184   | Abr32079<br>Abr32127<br>Adv31901<br>Adv31853 |
| Human 109 Human 109 Wheat gli Aspergill Aspergill Human SNP Human BLy Human BLY Human BLY Human B L Human B L Human B L Human B L Human B C Human |  |

## ALIGNMENTS

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RESULT 1
ADR10611
ID ADR10611
XX ADR1
AC ADR1
XX ADR1

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Horse IgE epitope recognised
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JAN-2004; 2004WO-US003566.
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Hammerberg B;

WPI; 2004-593545/57.

Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian Igor treating asthma or anaphylactic shock. IgE

Example 6; Page 9; 14pp; English.

The present invention relates to a novel monoclonal antibody (I) that specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. (I) is useful for testing an allergen reactivity of an IgE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut and corn allergens. The sample is a biological sample collected from a dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal antibodies recognise epitopes on canine IgE corresponding to amino acid residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the

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RESULT 2
ADC64569
ID CADC6
XX ADC6
XX ADC6
XX ADC6
XX ADC6
XX ADC6
XX HORE
XX CREC
XX 
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The invention relates to an immunogenic composition comprising an CC isolated polypeptide having an amino acid sequence that is at least 80% CC identical to 6 (S1-S6), 15 amino acid sequence that is at least 80% CC identical to 6 (S1-S6), 15 amino acid peptide sequences derived from CC equine immunoglobulin E (the composition induces production of an CC antibody that specifically binds to equine immunoglobulin (Ig)E), the six CC polypeptides are not explicitly identified in the specification. Also CC included are a composition comprising an antibody that specifically binds to equine IgE made by the process of immunising an CC animal with a polypeptide at least 80% identical to (S1)-(S6), an antibody that specifically binds to equine IgE (involving immunising an CC animal with a composition further comprising an isolated polypeptide (the CC amino acid sequence of the polypeptide is at least 80% identical to (S1)-(S6), and collecting antiserum from the animal) and a kit for detection of equine IgE in a biological sample comprising the antibody and means CC for detecting specific binding of the antibody to equine IgE. The CC antibody is useful for detecting equine IgE protein in a biological sample (serum) which involves contacting the sample with the antibody, the antibody and antigen are inspected of the antipen/antibody complex. The antibody and antigen are immobilised on a solid surface. The antibody is labelled such that the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  밁
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunogenic composition comprising an isolated equine immunoglobulin polypeptide that induces production of antibodies which specifically to equine immunoglobulin E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and 3.76 were observed to have good cross-reactivity with the epsilon-chain of IgE from cat and horse, but did not exhibit cross-reactivity with either pig or human epsilon-chains of IgE. The present sequence is the horse IgE 3.76 recognition site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI;
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Best Local
           This invention describes novel antibodies that immunospecifically bind to B Lymphocyte Stimulator (BLyS) polypeptides. BLyS is a member of the tumour necrosis factor (TNF) super family and induces B cell proliferation and differentiation. The antibodies of the invention have cytostatic, immunosuppressive, immunostimulant, immunomodulatory, antirheumatic and antiAIDS activity and can be used in vaccines to inhibit the expression and activity of BLyS. The antibodies bind to BLyS and so may be used to detect and quantitate the presence of BLyS in biological samples and may be used in this way to diagnose disease associated with aberrant expression of BLyS. They may also be administered to treat diseases associated with aberrant BLyS expression and activity such as cancer, immune, and autoimmune disorders and
                                                                                                                                                                                                                                                                                                                                                                                                         16-JUN-2000;
17-OCT-2000;
16-MAR-2001;
21-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BLys; B lymphocyte stimulator; TNF superfamily; human; cytostatic; tumour necrosis factor; B cell proliferation; B cell differentiation; immunosuppressive; immunostimulant; immunomodulatory; antirheumatic; antiAlDS; vaccine; cancer; immune; autoimmune disorder; immunodeficiency; systemic lupus erythematosus; rheumatoid arthritis; CVID; AlDS; common variable immunodeficiency; acquired immunodeficiency syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 complex can be detected. The complex is detected using a second labelled antibody. The peptides are useful for generating antibodies specific for IgE which can serve as a diagnostic test for allergy. The present sequence is a Horse immunoglobulin E, IgE, heavy chain immunogenic peptide from the middle portion of the C2 region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABP46438
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human BLyS binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200202641-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                  (CAMB-)
                                                                                                                                                                                                                                                                                                                                                                  (HUMA-)
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                                                                                                                                                                                                                                                                                                                                                                HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                  CAMBRIDGE ANTIBODY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                              Page 2992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IDGQKVDEQFPQHG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IDGQKVDEQFPQHG
                                                                                                                                                                                                                                                                                                                    Barash SC,
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2001US-0276248P.
2001US-0277379P.
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                                                                                                                                                                                                            3148pp; English.
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100.0%;
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                                                                                                                                                                                                                                          cancers and immune
                                                                                                                                                                                                                                                                                                                                                  TECHNOLOGY.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                   Vaughan
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2.3e-05;
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                                                                                                                                                                                                                                            polypeptides, useful 
ne disorders.
                                                                                                                                                                                                                                                                                                                      Hilbert
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e.g. systemic

lupus erythematosus,

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                                                    This invention relates to novel antibodies that immunospecifically bind to B lymphocyte stimulator (BLyS). The BLyS gene has been mapped to CC chromosome 13q34 and encodes a protein that is a member of the tumour CC necrosis factor superfamily and induces both in vivo and in vitro B cell proliferation and differentiation. Specifically, it refers to single CC chain antibody molecules (scFvs) derived, preferably, from the variable heavy CDR3 region that immunospecifically bind to a polypeptide, or CC fragment thereof, of either human, murine, rat or monkey BLyS. The present invention refers to the use of such antibodies in various methods CC for the detection, diagnosis and prognosis of diseases related to the aberrant expression or inappropriate function of BLyS or its receptor. As such, these compositions are useful for identifying immune disorders including myasthenia gravis and multiple sclerosis, inflammatory CC disorders e.g. asthma and rheumatoid arthritis, infectious diseases such as ALDS and proliferative disorders including leukaemia, carcinoma and CC lymphoma, Arcordingly rhew can be described as activities of the content of the conte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       B cell proliferation; differentiation; scFv; myasthenia gravis; multiple sclerosis; asthma; rheumatoid arthritis; AIDS; leukaemia; carcinoma; lymphoma; antirheumatic; antiarthritic; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunodeficiency (e.g. common variable immunodeficiency (CVID) and acquired immunodeficiency syndrome (AIDS)). ABP43990-ABP47228 represent the antibodies and fragments of the antibodies described in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 14 AA;
                lymphoma. Accordingly, they can be described as exhibiting various activities such as antirheumatic, antiarthritic, neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-505530/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ruben
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19-DEC-2001; 2001US-0340817P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antibody; B
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  antiinflammatory, antiasthmatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel antibody that immunospe
(BLys), useful for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                arthritis, asthma and
                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 2449;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       lymphocyte
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunospecifically binds to detecting and treating disease
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100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 36;
Pred. No.
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  antiallergic and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  leukemia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a B lymphocyte stimulator
ses or disorders e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BLyS SeqID
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ARESULT 5
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Best Local S
Matches 6
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17-OCT-2000;
16-MAR-2001;
21-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antiarthritic; Neuroprotective; Muscular-Gen.; Antiasthmatic; Antiallergic; Antimicrobial; Anti-HIV; Cytostatic; B-lymphocyte; antibody; autoimmune disease; b-cell lymphoma; systemic lupus erythematosus; rheumatoid arthritis; immune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 14
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                                                                                                                                                                                                    New antibodies that immunospecifically binds to B Lymphocyte Stimulator protein, useful for diagnosing, treating, or preventing autoimmune disease, e.g. systemic lupus erythematosus or rheumatoid arthritis, or l
                                                                                                                                                                                                                                                                                                                                                                                                                                        (RUBE/)
(BARA/)
(CHOI/)
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CHOI G H.
VAUGHAN T.
HILBERT D.
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BARASH S C.
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2001US-0880748.
2001US-0331469P.
2001US-0340817P.
2002US-00293418.
2002US-0293418.
2004US-0583296P.
2004US-0580347P.
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2001US-0276248P.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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1.1e+02;
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                                                                                                                                                                                                                                                                                                                                                  Hilbert
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The invention relates to a novel antibody that immunospecifically to a B Lymphocyte Stimulator protein. The protein comprises an and sequence that is 85% identical to the VH domain of any one of the chain variable fragments (gcFvs) of SEQ ID NOs. 1-2128, and/or an acid sequence that is at least 85% identical to the VL domain of a

Example 1; SEQ ID NO 2449; 240pp;

English

immunospecifically binds

w

single-amino

one

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RESULT 6
AEE33637
ID AEE33
XX AEE3
AC AEE3
XX Whea
XX Immu
KW T-ce
KW Gast
XX 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local
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                                                                                                                                                                                                                                                Preventing or treating celiac disease comprises administering to an individual a peptide (analogue) comprising at least one T cell epitope (e.g. a wheat and/or an oat epitope).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-APR-2004; 2004AU-00201774.
11-FEB-2005; 2005AU-00900650.
                                                                                                                                                                                 Claim 1;
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                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2005-769484/78.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AEE33637
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100.0%; Pred. No.
Live 0; Mismatc
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                                                                                                                                                                              94pp;
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                                                                                                                                                                              English.
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5. 1.1e+02;
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The invention relates to a method of preventing or treating celiac disease comprises administering to an individual at least one agent selected from a peptide comprising at least one epitope, given in the specification, or their equivalents, and an analogue of the peptide,

Query Match

36.5%;

Score

35;

83 9

Length

16;

Sequence

16

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RESULT 7
ARES34193
ID ARES3
AX ARES34193
XX ARES3
XX Whea
XX Immu
KW Immu
KW Immu
KW T-cee
KW Gast
XX HO20
XX 
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                                                    The invention relates to a method of preventing or treating celiac disease comprises administering to an individual at least one agent selected from a peptide comprising at least one epitope, given in the specification, or their equivalents, and an analogue of the peptide, which is capable of being recognized by a T cell receptor that recognizes the peptide. The methods, agents, antagonists and compositions are useful for treating or preventing celiac disease. The agent or antagonist, or wild type sequence is useful for producing an antibody specific to the agent, antagonist or wild type sequence. The mutation in an epitope of a gluten protein is useful for decreasing the ability of the gluten protein co cause celiac disease. The present sequence represents the amino acid sequence of a wheat gliadin/glutein epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               which is capable of being recognized by a T cell receptor that recognizes the peptide. The methods, agents, antagonists and compositions are useful for treating or preventing cellac disease. The agent or antagonist, or wild type sequence is useful for producing an antibody specific to the agent, antagonist or wild type sequence. The mutation in an epitope of a gluten protein is useful for decreasing the ability of the gluten protein to cause celiac disease. The present sequence represents the amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Preventing or treating celiac disease comprises administering to an individual a peptide (analogue) comprising at least one T cell epitope (e.g. a wheat and/or an oat epitope).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 879; 94pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-APR-2004;
11-FEB-2005;
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T-cell-receptor-antagonist; Vaccine; celiac disease; immunogenicity;
gastrointestinal disease; immune disorder; gluten protein; gliadin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wheat gliadin/glutein epitope SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Beissbath T,
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2005AU-00900650.
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54.5%;
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Pred. No. 1.8e+02;
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RESULT 8
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                                                                                                                                              The present invention relates to a novel monoclonal antibody (I) that CC specifically binds to a mammalian IgE peritope, where the epitope is CC between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. CC (I) is useful for testing an allergen reactivity of an IgE sample. The cand corn allergens. The sample is a biological sample collected from a CC and corn allergens. The sample is a biological sample collected from a CC dog, cat or horse. (I) is also useful for detecting mammalian IgE and for CC treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal CC antibodies recognise epitopes on canine IgE corresponding to amino acid CC residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the CC canine IgE speilon-chain. Recognition of epsilon-chains from IgE from CC cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and CC of IgE from cat and horse, but did not exhibit cross-reactivity with the cepsilon-chain epsilon-chains of IgE. The present sequence is the CC cat IgE 3.76 recognition site.
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Matches 6; Conserv
                                                          Matches
                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian IgE or treating asthma or anaphylactic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody;
                                                                                                                    Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                    Example 6; Page 9; 14pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cat IgE epitope recognised by monoclonal antibody 3.76,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-OCT-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADR10610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hammerberg
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                                                                       ocal
                                                        Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OKVDEQFPOHG 14
                           IDGOKVDEQFPQHGLVKQ 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     QQPEQQFPQPG 16
VDGQKATNI FPYTAPGKQ
                                                                                                                      $
                                                          Conservative
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                                                                       36.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               54.5%;
18
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                                                                       Score 35; DB 8;
Pred. No. 2e+02;
                                                        Mismatches
                                                          9; Indels
                                                                                       Length 18;
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Best Local S
Matches 5
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25-MAR-2003
09-OCT-1994
                                                                                                                                                                                                                                                                                                                                                  The surface M protein of Group A streptococi is the major virulence factor and protective antigen of these organisms. However, there are a tremendous number of M protein serotypes. The invention provides recombinant M protein antigens comprising a gene encoding a carrier protein and an NH2 or COOH terminal M protein fragment carrying one or more epitopes. AAR50332 is a suitable NH2-terminal fragment of M3 protein for constructing antigens which elicit opsonic antibodies in an immunised animal when linked or fusied to an appropriate carrier. (Updated on 25-MAR.2003 to correct PN field.) (Updated on 27-AUG-2003 to correct OS
          Human; 121P2A3; cytostatic; immunostimulant; vaccine; SSH; humoral immune response; cellular immune response; suppression subtractive hybridisation; HLA; human leukocyt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B subunit; labile toxin; M protein; fur Group A streptococci; rheumatic fever;
                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 11; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New recombinant hybrid streptococcal M protein antigen(s) - which elicit opsonic antibodies without eliciting cross-reactive antibodies to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence of
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                                                           HLA protein 121P2A3 peptide #11048.
                                                                                        28-MAR-2003
                                                                                                                ABP94693;
                                                                                                                                          ABP94693 standard; peptide; 15 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mammalian heart tissue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1994-118162/14
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                                                                                                                                                                                                                                                                                                                                        field.)
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                                                                                                                                                                                                                                 2 DGQKVDEQFPQH 13
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                                                                                                                                                                                                                                                           Similarity 41. 5; Conservative
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(revised)
(first entry)
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                                                                                                                                                                                                                                                                       35.4%;
41.7%;
                                                                                      entry)
                                                                                                                                                                                                                                                           Score 34; DB 2; Le
Pred. No. 2.4e+02;
4; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fusion protein;
er; pharyngitis;
              human leukocyte antigen.
                                                                                                                                                                                                                                                                                    Length 15;
                                                                                                                                                                                                                                                             Indels
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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25-APR-2001; 2001US-0286630P
22-JUN-2001; 2001US-0300373P
                                                                                suppression
                                                                                                                  Human; 121P2A3; cytostatic; immunostimulant; vaccine; SSH;
                                                                                                                                                   HLA protein 121P2A3 peptide
                                                                                                                                                                                                                                                     ABP95588 standard;
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Afar DEH, Saffran
                   WO200283068-A2
                                                 Homo sapiens.
                                                                                                    humoral
                                                                                                                                                                                      28-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 6; Conserv
                                                                                immune response; cellular immune response;
sion subtractive hybridisation; HLA; human
                                                                                                                                                                                                                                                                                                                                                                       DGQKVDEQFPQHGLV 16
                                                                                                                                                                                                                                                                                                                                        ENEKLDRQHVQHQLL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Raitano AB, I
                                                                                                                                                                                                                                                       peptide;
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Pred. No. 2.4e+02;
Pred. marches 5; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hubert RS,
ison RK, Ge
                                                                                  leukocyte
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Best Local
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25-APR-2001;
22-JUN-2001;
                                                                                                                                                                                                                                                                Human; 121P2A3; cytostatic; immunostimulant; vaccine; SSH; humoral immune response; cellular immune response; suppression subtractive hybridisation; HLA; human leukocyte antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABP95194 standard;
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                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                     HLA protein 121P2A3 peptide #11549
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121P2A3 polypeptides, useful for eliciting humoral or cellular immune
responses or in assessing the status of 121P2A3 gene products in normal
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Afar DEH, Saffran D, Morrisc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   represent peptides from the 121P2A3 variants of the invention
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2001US-0286630P.
2001US-0300373P.
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Pred. No. 2.4e
4; Mismatches
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n K, M
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2.4e+02;
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ison RK,
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W, Jakob
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09-APR-2002; 2002WO-US011359.

24-OCT-2002.

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Best Local Similarity
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Afar DEH, Saffran D,
                                                                                                                                                                                                            tissue Factor VIIa; tFVIIa; chronic thromboembolic disease; fibrin formation; vascular disorders; deep venous thrombosis; arterial thrombosis; stroke; atherosclerosis; septicaemia.
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25-APR-2001; 2001US-0286630P.
22-JUN-2001; 2001US-0300373P.
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                                              06-FEB-2002;
                                                                               30-JAN-2003; 2003US-00356257.
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               (GETH ) GENENTECH INC
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                                              2002US-0355420P
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                                                                                                                                                                                                                                                                             (tfVIIa)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 35.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Morrison
                                                                                                                                                                                                                                                                             peptide antagonist
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 34; DB 6;
Pred. No. 2.4e+02;
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Faris M,
on K, Mori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      s M, Hubert RS, Mitchell Morrison RK, Ge W, Jakob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5
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RESULT 14
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The present invention relates to a novel monoclonal antibody (I) that specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE (I) is useful for testing an allergen reactivity of an IgE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut and corn allergens. The sample is a biological sample collected from a
                                                                                                                                                                                                                                                                                Novel antibody that specifically binds to for testing an allergen reactivity of IgE or treating asthma or anaphylactic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide, useful for preventing or treating chronic thromboembolic diseases or disorders associated with fibrin formation including vascular disorders, such as deep venous thrombosis, arterial thrombosis, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17 AA;
                                                                                                                                                                                                                              Example 6; Page 9; 14pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-593545/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-AUG-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADR10609 standard; peptide; 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-JAN-2003; 2003US-0440472P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JAN-2004; 2004WO-US003566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IgE epitope recognised by monoclonal antibody 3.76,
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Pred. No. 2.7e+02;
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                                                                                                                                                                                                                                                                                                                        mammalian IgE epitope, useful sample, detecting mammalian I
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                                                                                                                                                                                                                                                                                                                               mammalian IgE
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RESULT 15
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                                                      The present invention describes an isolated DNA (I) comprising a nucleotide sequence coding for an enzyme having enone reductase activity. Also described: (I) a vector or a plasmid comprising (I); (2) a host cell transformed or transfected by (I) or the vector or the plasmid of (I); (3) a polypeptide encoded by (I) and (4) a process for the production of levodione which comprises contacting ketoisophorone with the polypeptide of (3) or with the host cell of (2) or a cell-free extract under conditions for the production of levodione, e.g. at pH 4.0-9.0 and at a temperature of 10-60 degrees Celsius for 5 minutes to 72 hours or at pH 5.0-8.0 and at a temperature of 20-60 degrees Celsius for 15 minutes to 48 hours. The DNA (I) and the encoded polypeptide can be used in improving the production process of condition, which is important in the synthesis of optically active carotemoids. The present sequence represents a Candida kefyr (Kluyyeromyces marxianus) enone reductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal antibodies recognise epitopes on canine IgE corresponding to amino acid residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from cat, horse, pig and human by 5.91 and 3.76 were observed to have good cross-reactivity with the epsilon-chain of IgE from cat and horse, but did not exhibit cross-reactivity with either pig or human epsilon-chains of IgE.
                                                                                                                                                                                                                                                                                                                                                                                                                            New enone reductase, useful in improving the production process of levodione, which is important in the synthesis of optically active
    Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                             Example 1; SEQ ID NO 9; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              enone reductase; enzyme; levodione; ketoisophorone; carotenoid; Candida kefyr; Kluyveromyces marxianus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-295422/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kataoka M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-SEP-2002; 2002EP-00021098
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Candida kefyr enone reductase peptide SEQ ID NO:9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADL98126;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kluyveromyces marxianus
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                                         which
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                                     the exemplification of the present invention.
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1; Mismatches
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2.9e+02;
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71.4%;
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                                                Score 33; DB 8;
Pred. No. 2.2e+02;
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Listing first 45 summaries
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1: /RMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /RMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US09_PUBCOMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US11_PUBCOMB.pep:*
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US-09-800-748-2449
US-11-054-515-2449
US-11-054-515-2449
US-11-054-65-10
US-10-758-165-11
US-10-758-165-7-11
US-10-758-165-9
US-11-127-804-9
US-11-127-804-9
US-11-127-804-9
US-11-079-709-3
US-11-086-478-842
US-11-086-478-842
US-11-086-478-842
US-11-266-478-215-2703
US-09-880-748-215-216-219-418-2151
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     11 Appli
2449, Ap
21449, Ap
2449, Ap
251, Appli
252, Appli
273, Appli
2446, Ap
2703, Ap
2703, Ap
2703, Ap
2704, Ap
2704,
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| RESULT 2 US-10-052-788-5 ; Sequence 5, Application US/10052788 ; Publication No. US20030087314A1 ; GENERAL INFORMATION: APPLICANT: Gershwin, Laurel J. APPLICANT: Marina, Warren V. APPLICANT: The Regents of the University of California TITLE OF INVENTION: Epsilon Immunoglobulin Chain Derived TITLE OF INVENTION: Induction of Anti-IgE Antibodies FILE REFRENCE: 023070-121000US CURRENT APPLICATION NUMBER: US/10/052,788 CURRENT PILING DATE: 2001-11-08 NUMBER OF SEQ ID NOS: 6 SOFTWARE: Patentin Ver. 2.1 SEQ ID NO 5 LENGTH: 15 TYPE: PTI ORGANISM: Artificial Sequence | Query Match  Query Match  Best Local Similarity 100.0%; Pred. No. 9.4e-09;  Matches 18; Conservative 0; Mismatches 0;  Qy 1 IDGQKVDEQFPQHGLVKQ 18 | RESULT 1  US-10-758-165-11  US-10-758-165-11  Sequence 11, Application US/10758165  Publication No. US20050196816A1  GENERAL INFORMATION:  APPLICANT: Hammerberg, Bruce  TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MITTLE REFERENCE: 5051-661  CURRENT APPLICATION NUMBER: US/10/758,165  CURRENT FILING DATE: 2004-01-16  PRIOR APPLICATION NUMBER: US 60/440,472  PRIOR FILING DATE: 2003-01-16  NUMBER OF SEQ ID NOS: 16  SOUTWARE: PatentIn version 3.2  SEQ ID NO 11  LENGTH: 18  TYDE: PRT  ORGANISM: Equus caballus  US-10-758-165-11 | ALIGNMENTS | 28 31 32.3 14 6 US-11-054-915-2151 29 31 32.3 14 6 US-11-266-444-2146 30 31 32.3 14 6 US-11-266-444-2151 31 31 32.3 15 4 US-10-059-261-108 32 31 32.3 15 4 US-10-059-261-216 33 31 32.3 15 5 US-10-627-649-108 34 31 32.3 15 5 US-10-627-649-216 35 31 32.3 18 5 US-10-758-165-15 36 30 31.2 10 4 US-10-353-929-166 37 30 31.2 11 3 US-09-977-827-8 38 30 31.2 17 3 US-09-977-827-13 40 29 30.2 11 3 US-09-977-827-13 41 29 30.2 11 3 US-09-977-827-13 42 29 30.2 11 3 US-09-978-27-13 43 29 30.2 11 5 US-10-500-878-14 44 29 30.2 13 6 US-11-52-974A-272 43 29 30.2 13 6 US-11-52-974A-272 44 29 30.2 13 6 US-11-52-974A-272 45 29 30.2 15 4 US-10-282-960-23 |
|--|---|--|------------|--|
| nia<br>Ved Peptides for  | Length 18;<br>Indels 0; Gaps 0;   | IN MAMMALIAN SPECIES   |            | Sequence 2151, Ap<br>Sequence 2146, Ap<br>Sequence 2151, Ap<br>Sequence 216, App<br>Sequence 216, App<br>Sequence 216, App<br>Sequence 166, App<br>Sequence 15, Appli<br>Sequence 4580, A<br>Sequence 314, App<br>Sequence 13, Appli<br>Sequence 13, Appli<br>Sequence 272, App<br>Sequence 272, App<br>Sequence 272, App<br>Sequence 273, Appli<br>Sequence 273, Appli  |

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FILE REFERENCE: PF523P2
CURRENT APPLICATION NUMBER: US/10/293,418
CURRENT FILING DATE: 2002-11-27
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILING DATE: 2001-12-19
PRIOR PRIOR PLING DATE: 2001-06-15
PRIOR PRIOR DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR APPLICATION NUMBER: 60/297,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/276,248
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US-09-880-748-2449
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PRIOR FILING DATE: 2000-06-15
PRIOR PRIOR PRILING DATE: 2000-10-17
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 3239
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US-09-880-748-2449
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Publication No. US20030223998A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 2449
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
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CURRENT FILING DATE: 2001-06-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Ruben et al. TITLE OF INVENTION: Antibodies that Immunospecifically Bind FILE REFERENCE: PF523
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2001-03-16
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100.0%; Pr
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100.0%; Pr/
100.0%; 0;
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Pred. No.
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63;
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7.4e-06;
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Sequence 2449, Application US/11266444
Publication No. US20060062789A1
GENERAL INFORMATION:
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                                                                                                                           ; ORGANISM: Homo US-11-054-515-2449
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-293-418-2449
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US-11-054-515-2449
                                                           Query Match
Best Local S
Matches 6
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PRIOR FILING DATE: 2004-02-11
PRIOR APPLICATION NUMBER: 60/580,347
PRIOR FILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
                                                                                                                                                                        SEQ ID NO 2449
LENGTH: 14
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SEQ ID NO 2449
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Remaining Prior Application d
NUMBER OF SEQ ID NOS: 3247
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CURRENT FILING DATE: 2005-02-10
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TITLE OF INVENTION:
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PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-16
                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 2001-06
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                                                          Local Similarity
nes 6; Conserv
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                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/277,379
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                             FPQHGL 15
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                                                             Conservative
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                                                                          100.0%;
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                                                                            Score 36;
Pred. No.
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Pred. No.
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                                                                          . 63;
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63;
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APPLICANT: Ruben et al

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US-10-141-627-11
Sequence 11, Application US/10141627
Publication No. US20020176863A1
GENERAL INFORMATION:
APPLICANT: Dale, James B.
TITLE OF INVENTION: ANTIGEN OF HYBRID M PROTEIN AND CARRIER TITLE OF INVENTION: FOR GROUP A STREPTOCOCCAL VACCINE
FILE REFERENCE: 481112.404C3
CURRENT APPLICATION UNMER: US/10/141,627
CURRENT FILING DATE: 2002-05-07
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US-10-758-165-10
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US-10-758-165-10
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US-11-266-444-2449
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CURRENT APPLICATION NUMBER: US/11/266,444

CURRENT FILING DATE: 2005-11-04

PRIOR APPLICATION NUMBER: 09/880,746

PRIOR FILING DATE: 2001-06-15

PRIOR APPLICATION NUMBER: 60/212,210

PRIOR APPLICATION NUMBER: 60/214,210

PRIOR FILING DATE: 2000-06-16

PRIOR APPLICATION NUMBER: 60/240,816

PRIOR APPLICATION NUMBER: 60/240,816

PRIOR APPLICATION NUMBER: 60/240,816
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Publication No. US20050196816A1
GENERAL INFORMATION:
APPLICANT: Hammerberg, Bruce
TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION
ETT TO THE TOTAL TOT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn version 3.2
SEQ ID NO 10
LENGTH: 18
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PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
NUMBER OF TAKE 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR FILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
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NUMBER OF SEQ ID NOS: 3239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
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llarity 44.4%;
Conservative
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; Pred. No. 1.2e
1; Mismatches
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OTHER INFORMATION: NH2-terminal fragment of M protein for OTHER INFORMATION: constructing antigens, which elicit opsonic; OTHER INFORMATION: antibodies in an immunized animal US-10-141-627-11
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US-10-356-257-255
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                                  US-10-758-165-9
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US-10-758-165-9
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SEQ ID NO 11
LENGTH: 15
                                                                                                      APPLICANT: Hammerberg, Bruce
TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR FILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn version 3.2
SEQ ID NO 9
                                                                                                                                                                                                                                                                                        Sequence 9, Application US/10758165
Publication No. US20050196816A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/355,420
PRIOR ELLING DATE: 2002-02-06
NUMBER OF SEQ ID NOS: 355
SEQ ID NO 255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: LAZARUS, ROBERT A.
APPLICANT: MAUN, HENRY R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 255, Appublication No.
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Best Local Similarity
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: FVIIA Antagonists
FILE REFERENCE: P1950R1
CURRENT APPLICATION NUMBER: US/10/356,257
CURRENT FILING DATE: 2003-01-30
                                                LENGTH: 18
TYPE: PRT
ORGANISM: Canis familiaris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE: OTHER INFORMATION: sequence is synthesized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 17
TYPE: PRT
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                              1 GEGVEEEFPE 10
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41.7%;
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 Mismatches

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Pred. No. 1.7e+02;
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Length 18;
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Query Match

35.4%;

Score 34;

DB

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APPLICANT: Lin, Rong-Hwa
APPLICANT: Chang, Chung Nan
TITLE OF INVENTION: T-CELL DEATH-INDUCING EPITOPES
FILE REFERENCE: 113062-009001
CURRENT APPLICATION NUMBER: US/11/127,804
CURRENT FILING DATE: 2005-05-11
PRIOR APPLICATION NUMBER: US 60/570,161
PRIOR APPLICATION NUMBER: US 60/570,161
PRIOR FILING DATE: 2004-05-11
NUMBER OF SEQ ID NOS: 23
SOSTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 9
LENGTH: 13
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US-10-079-709-3
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Best Local Similarity
Watches 6; Conserve
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US-11-127-804-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 3, Application US/10079709 Publication No. US20030119163A1
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Publication No. US20060003940A1
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Matches 6; Conservative
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                              SOFTWARE:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 02-FEB-200:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                     ZIP: 94025-3471
COMPUTER READABLE FORM:
EDDIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster
STREET: 545 Middlefield Road, Suite 200
CITY: Menlo Park
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                  COUNTRY:
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                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                                                                                                Gerardus Selten
/ENTION: Cloning and Expression of Microbial
                                                                                                                                                                                                                                                                                                                                                                                                                  Annemarie E. Veenstra Rudolf G.M. Luttin
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Robert F.M. Van Gorcom
Willem Van Hartingsveldt
Petrus A. Van Paridon
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                                                                                                                         PatentIn Release #1.0,
                                                     02-FEB-2002
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                                                                    US/10/079,709
09/233,510
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Pred. No. 1.8e+02;
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1; Mismatches 4
                                                                                                                           Version #1.25
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US-11-036-272-3
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                     Sequence 3, Application US/11036272 Publication No. US20060063243A1
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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TELEFAX: 415-327-2951
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
APPLICANT:
REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-327-7250
                                                                                                                                                                                                   SOFTWARE:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/036,272
FILING DATE: 14-JAN-2005
                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide
HYPOTHETICAL: NO
FRAGMENT TYPE: N-terminal
                                                                                                                                                                      CLASSIFICATION:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 20-JAN-1999
APPLICATION NUMBER: 07/6:
FILING DATE: 24-MAY-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                            STREET: 545 Middlefield Road, Suite 200
CITY: Menlo Park
STATE: California
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Local Similarity 66.7%;
les 6; Conservative
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REFERENCE/DOCKET NUMBER: 24
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Willem Van Hartingsveldt
Petrus A. Van Paridon
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                                                                                                                                                                                                                                                                     PatentIn Release #1.0, Version #1.25
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Pred. No. 2e+02;
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INFORMATION FOR SEQ ID NO:

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Sequence 2703, Application US/09880748

Publication No. US20030059937A1

GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523

CURRENT APPLICATION NUMBER: US/09/880,748

CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-15
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION UMMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
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PRIOR FILING DATE: 2000-06-15
PRIOR PRILING DATE: 2000-06-15
PRIOR PRILING DATE: 2000-10-17
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-17
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
PRIOR FILING DATE: 2001-05-25
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Publication No. US20030059937A1

GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523
CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-66-15
DETERM TOTATION TOTATION TOTATION NUMBER: US/09/880,748
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Best Local Similarity 66.7%;
Matches 6; Conservative
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SEQ ID NO 2446
LENGTH: 14
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LENGTH: 10 amino acids
TYPE: amino acid
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                    US-11-32-378-136

US-11-140-487A-2148

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US-11-244-348A-7

US-11-32-356-9

US-11-32-378-216

US-11-140-487A-795

US-11-140-487A-795

US-11-140-487A-1029

US-11-122-986-542

US-11-122-986-673

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US-11-23-866-79

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Sequence 2148, Ap
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|  | -11-     | -11      | US-11-140-487A-1506 | US-11-219-563-16 | -10-538-066 | -066 | -10-538-066-32 | -10-538 | 3-10-538-066-13 | -11 | 11       | -11     | )-11-    | US-11-054-072-89 | -10-538-066- | -10-538-066-32 | -10-546-594-3 | -11-122-986-66 | -11-122-986-39 | T18-924- |
|  | 60, Appl | 622, App | 1506, Ap            | 16, Appl         | 394, App    | -    | •              | •       | _               |     | 59, Appl |         | 1608, Ap | 89, Appl         | 346, App     | 326, App       | 38, Appi      | 669, App       | 398, App       | Trddw 'F |

### ALIGNMENTS

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RESULT 1
US-11-332-378-136
; Sequence 136, Applicati
; Publication No. US2006(
; Publication No. US2006(
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; Publication No. US2006(
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                                                                                                                                           Sequence 2148, Application US/11140487A Publication No. US20060093617A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/NO99/00220
PRIOR FILING DATE: 1999-06-30
NUMBER OF SEQ ID NOS: 231
SOFTWARE: PATENTIN VERSION 3.0
SEQ ID NO 136
                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Saeboe-Larsen, Stein
TITLE OF INVENTION: Antigenic Peptides Derived from Telomerase
FILE REFERENCE: 01702.401800
CURRENT APPLICATION NUMBER: US/11/332,378
CURRENT FILING DATE: 2006-01-17
PRIOR APPLICATION NUMBER: US/09/743,281
PRIOR FILING DATE: 2001-07-05
APPLICANT: Innogenetics N.V.
TITLE OF INVENTION: Peptides for inducing a
FILE REFERENCE: 166
CURRENT APPLICATION NUMBER: US/11/140,487A
CURRENT FILING DATE: 2005-05-31
PRIOR APPLICATION NUMBER: EP 04012951.2
PRIOR FILING DATE: 2004-06-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Gaudernack, Gustav
APPLICANT: Erikeen, Jon Amund
APPLICANT: Moller, Mona
APPLICANT: Gjertsen, Marianne Klemp
APPLICANT: Saeterdal, Ingvil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 9
TYPE: PRT
ORGANISM: Homo sapiens, other or unknown
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5.2e+04;
                                                                                                          CTL and/or HTL
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CURRENT APPLICATION NUMBER: US/10/544,499
CURRENT FILING DATE: 2005-08-03
PRIOR APPLICATION NUMBER: 60/447,590
PRIOR FILING DATE: 2003-02-14
PRIOR FILING DATE: 2004-02-10
RIOR FILING DATE: 2004-02-10
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 3
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetic peptide mimotope
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PRIOR FILING DATE: 2004-10-28
PRIOR FPLICATION NUMBER: EP 05102441.2
PRIOR FILING DATE: 2005-03-25
PRIOR PILING DATE: 2004-06-03
PRIOR FILING DATE: 2004-06-03
PRIOR PILING DATE: 2004-10-29
PRIOR FILING DATE: 2004-10-29
PRIOR PILING DATE: 2004-10-29
PRIOR APPLICATION NUMBER: US 60/65,395
PRIOR APPLICATION NUMBER: US 60/65,395
PRIOR PILING DATE: 2005-03-25
NUMBER OF SEQ ID NOS: 2278
SOFTWARE: PATENTIN VETBION 3.1
SEQ ID NO 2148
                                                                                                                                                                              US-11-281-644A-2
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US-11-140-487A-2148
                                                                                                                  Sequence 2, Application US/11281644A Publication No. US20060105953A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
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              APPLICANT: Arnaud Lacoste
APPLICANT: Christopher J. Evans
TITLE OF INVENTION: SYSTEM FOR PROTEASE MEDIATED
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: 30435-164-US-U1
  CURRENT APPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Anthony J. Conley
APPLICANT: Beverly H. Galinski
APPLICANT: Allison Montalvo
TITLE OF INVENTION: Immunogens f
FILE REFERENCE: 21219P
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TYPE: PRT
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Local Similarity 50.0%;
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Local Similarity 46.2%;
les 6; Conservation
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NUMBER: US/11/281,644A
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Pred. No. 40;
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Pred. No. 27;
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US-11-122-986-795
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/423,875
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,832
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,915
PRIOR APPLICATION NUMBER: 60/423,757
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,757
PRIOR APPLICATION NUMBER: 60/423,758
PRIOR APPLICATION NUMBER: 60/423,758
                                                                                                                                                                                                     Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 844
SOFTWARE: PatentIn Ver. 3.3
SEQ ID NO 795
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PRIOR APPLICATION NUMBER: 60/628,694
PRIOR FILING DATE: 2004-11-17
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 2
                                                                   Matches
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                                                                                 Best Local Similarity
                                                                                                 Query Match
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Best Local
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PRIOR FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,373
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CURRENT APPLICATION NUMBER: US/11/122,986
CURRENT FILING DATE: 2005-05-05
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APPLICANT: DHARAMS
                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/424,367
PRIOR FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,376
PRIOR FILING DATE: 2002-11-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: VEDADI, MASOUD
TITLE OF INVENTION: ESSENTIAL NOVEL BACTERIAL
                                                                                                                                                                        LENGTH: 10
TYPE: PRT
                                                                                                                                                                                                                                                                              PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/424,370 PRIOR FILING DATE: 2002-11-06
                                                                                                                                                  ORGANISM: Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 16
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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FPEVGLKK 10
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57.1%;
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                                                                                   Score 25;
Pred. No.
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Pred. No. 93;
                                                                    Mismatches
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                                                                                   DB 81;
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US-11-301-554-2108

Sequence 2108, Application US/11301554

Publication No. US20060088527A1

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RESULT 7
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US-11-301-554-2108
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CURRENT APPLICATION NUMBER: US/11/301,554
CURRENT FILING DATE: 2005-12-13
PRIOR APPLICATION NUMBER: US 10/283,017
PRIOR FILING DATE: 2002-10-28
PRIOR APPLICATION NUMBER: US 10/113,872
PRIOR FILING DATE: 2002-03-28
PRIOR APPLICATION NUMBER: US 10/117,754
                                                                                                                                                                                 Sequence 9, Application US/10522356 Publication No. US20060105323A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 2157
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2108
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APPLICANT:
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FILE REFERENCE: 102286.155 US1
CURRENT APPLICATION NUMBER: US/10/522,356
CURRENT FILING DATE: 2005-01-26
PRIOR APPLICATION NUMBER: PCT/GB03/003192
                                                                                       APPLICANT: WHITELAW, CHRISTOPHER BRUCE ALEXANDER
APPLICANT: CLARK, ANTHONY JOHN
APPLICANT: WOLF, CHARLES ROLAND
TITLE OF INVENTION: MULTI-REPORTER GENE MODEL FOR TOXICOLOGICAL SCREENING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: MCNADD, Andria
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OR APPLICATION NUMBER: US 09/736,457
OR FILING DATE: 2000-12-13
OR FILING DATE: 2000-10-30
OR FILING DATE: 2000-10-30
OR APPLICATION NUMBER: US 09/677,419
OR APPLICATION NUMBER: US 09/677,419
OR FILING DATE: 2000-10-06
OR APPLICATION NUMBER: US 09/671,325
OR APPLICATION NUMBER: US 09/658,824
OR APPLICATION NUMBER: US 09/658,824
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FILING DATE: 2001-10-29
APPLICATION NUMBER: US 09/902,941
FILING DATE: 2001-07-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 2000-09-08
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5; Conserva
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Vedvick, Thomas S.
Bangur, Chaitanya S.
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45.5%;
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Pred. No. 1.2e+02
3; Mismatches
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PRIOR APPLICATION NUMBER: GB 0217402.7
PRIOR FILING DATE: 2002-07-26
NUMBER OF SEQ ID NOS: 41
SOFTWARE: PatentIn version 3.3
SEQ ID NO 9
LENGTH: 16
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TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR IN-
TITLE OF INVENTION: AGAINST PSEUDOMONAS INFECTION
FILE REFERENCE: 10901-0.15-999
CURRENT APPLICATION NUMBER: US/11/244,348A
CURRENT FILING DATE: 2005-10-04
PRIOR APPLICATION NUMBER: 60/616,125
PRIOR FILING DATE: 2004-10-04
NUMBER OF THE TOTAL TOTAL PRIOR PRIOR FILING DATE: 2004-10-04
NUMBER OF THE TOTAL TOTAL PRIOR PRIOR FILING DATE: 2004-10-04
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Best Local Similarity
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                                                                                                                                                                                                     APPLICANT: Gaudernack, Gustav
APPLICANT: Eriksen, Jon Amund
APPLICANT: Moller, Mona
APPLICANT: Gjertsen, Marianne
APPLICANT: Saeterdal, Ingvil
                                                          APPLICANT: Saeboe-Larsen, Stein
TITLE OF INVENTION: Antigenic Peptides Derived from
FILE REFERENCE: 01702.401800
CURRENT APPLICATION NUMBER: US/11/332,378
CURRENT FILING DATE: 2006-01-17
PRIOR APPLICATION NUMBER: US/99/743,281
PRIOR FILING DATE: 2001-07-05
PRIOR FILING DATE: 2001-07-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 12
SOFTWARE: FastSEQ for Windows Version
PRIOR APPLICATION NUMBER: PCT/NO99/00220
PRIOR FILING DATE: 1999-06-30
NUMBER OF SEQ ID NOS: 231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Pseudomonas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Unknown Sequence: ARF OTHER INFORMATION: epitope from unknown organism
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Saeterdal, Ingvil
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Moller, Mona
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66.7%;
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Pred. No. 1.4e+02;
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Pred. No. 1.5e+02;
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PRIOR APPLICATION NUMBER: EP 04012951.2

PRIOR APPLICATION NUMBER: EP 04012951.2

PRIOR FILING DATE: 2004-06-01

PRIOR APPLICATION NUMBER: EP 04447239.7

PRIOR FILING DATE: 2004-10-28

PRIOR APPLICATION NUMBER: EP 05102441.2

PRIOR FILING DATE: 2005-03-25

PRIOR APPLICATION NUMBER: US 60/576,310

PRIOR APPLICATION NUMBER: US 60/576,310
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TITLE OF INVENTION: Peptides for inducing a C
FILE REFERENCE: 166
CURRENT APPLICATION NUMBER: US/11/140,487A
CURRENT FILING DATE: 2005-05-31
PRIOR APPLICATION NUMBER: EP 04012951.2
PRIOR FILING DATE: 2004-06-01
PRIOR PHILING DATE: 2004-06-01
PRIOR PHILING DATE: 2004-05-03-25
PRIOR FILING DATE: 2004-05-03-25
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US-11-332-378-216
                                                                                                                                                                                                                                                                          Sequence 795, Applic Publication No. US20 GENERAL INFORMATION:
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Best Local
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                                                                                                                                                                                                              APPLICANT: Innogenetics N.V.
TITLE OF INVENTION: Peptides for inducing a CTL and/or HTL response to Hepatitis C
FILE REFERENCE: 166
                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/11/140,487A
CURRENT FILING DATE: 2005-05-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2005-03-25
NUMBER OF SEQ ID NOS: 2278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: hepatitis C virus
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Local Similarity 62.5%;
les 5; Conservation
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OR FILING DATE: 2005-03-25
OR APPLICATION NUMBER: US 60/576,310
OR FILING DATE: 2004-06-03
OR APPLICATION NUMBER: US 60/622,782
OR APPLICATION NUMBER: US 60/665,395
OR APPLICATION NUMBER: US 60/665,395
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Pred. No. 5.2e+04;
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US-11-122-986-542; Sequence 542, Application US/11122986; Publication No. US20060104989A1; GENERAL INFORMATION:
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US-11-140-487A-1029
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; TYPE: PRT
; ORGANISM: hepatitis C virus
US-11-140-487A-795
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PRIOR FILING DATE: 2005-03-25
NUMBER OF SEQ ID NOS: 2278
SOFTWARE: Patentin version 3.1
SEQ ID NO 755
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Best Local
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APPLICANT: EDWARDS, ALED
APPLICANT: DHARAMSI, AKIL
APPLICANT: VEDADI, MASQUD
TITLE OF INVENTION: ESSENTIAL NOVEL BACTERIAL POLYPEPTIDES
FILE REFERENCE: IPT-330.01
CURRENT FILLING DATE: 2005-05-05
PRIOR APPLICATION NUMBER: 60/423,875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 2278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/576,310
PRIOR FILING DATE: 2004-06-03
PRIOR APPLICATION NUMBER: US 60/622,782
PRIOR FILING DATE: 2004-10-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 166
CURRENT APPLICATION NUMBER: US/11/140,487A
CURRENT FILING DATE: 2005-05-31
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TITLE OF INVENTION: Peptides for inducing a CTL and/or HTL response to Hepatitis C
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62.5%;
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Pred. No.
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Pred. No. 1.8e+02;
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FILING DATE: 2002-11-05 APPLICATION NUMBER: 60/423,832 FILING DATE: 2002-11-05

FILING DATE: 2002-11-05

APPLICATION NUMBER: 60/423,915

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US-11-122-986-789
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                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/423,832
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,915
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,757
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,758
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Remaining Prior Application data removed - See File Wrapper NUMBER OF SEQ ID NOS: 844
SOFTWARE: PatentIn Ver. 3.3
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SOFTWARE: PatentIn Ver. 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: IPT-330.01
CURRENT APPLICATION NUMBER: US/11/122,986
CURRENT FILING DATE: 2005-05-05
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APPLICANT: DHARAMSI, AKIL
APPLICANT: VEDADI, MASOUD
TITLE OF INVENTION: ESSENTIAL NOVEL BACTERIAL POLYPEPTIDES
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                                                                OR FILING DATE: 2002-11-05
OR APPLICATION NUMBER: 60/424,367
OR FILING DATE: 2002-11-06
OR PILING DATE: 2002-11-06
OR PILING DATE: 2002-11-06
OR APPLICATION NUMBER: 60/424,370
OR FILING DATE: 2002-11-06
OR APPLICATION NUMBER: 60/424,362
OR APPLICATION NUMBER: 60/424,362
OR APPLICATION NUMBER: 60/424,363
OR APPLICATION NUMBER: 60/424,373
OR FILING DATE: 2002-11-06
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Local Similarity 30.0%;
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OR FILING DATE: 2002-11-06
OR FILING DATE: 2002-11-06
OR FILING DATE: 2002-11-06
OR APPLICATION NUMBER: 60/424,362
OR FILING DATE: 2002-11-06
OR APPLICATION NUMBER: 60/424,362
OR APPLICATION NUMBER: 60/424,373
OR FILING DATE: 2002-11-06
OR FILING DATE: 2002-11-06
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FILING DATE: 2002-11-05
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APPLICATION NUMBER: 60/424,367
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D. US20060104989A1
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Pred. No. 2.6e+02;
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; TYPE: PRT ; ORGANISM: Streptococcus pneumoniae US-11-122-986-789
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Search completed: May 30, 2006, 15:20:31 Job time : 4 secs
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                                                                                                                                                                                                                         ; LENGTH: 11,
TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-11-122-986-673
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US-11-122-986-673
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Best Local Similarity 66...
""" tohes 4; Conservative
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                                                                                                                                                 Matches
                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                 Remaining Prior Application data removed NUMBER OF SEQ ID NOS: 844 SOFTWARE: PatentIn Ver. 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/423,875
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,832
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,915
PRIOR FILING DATE: 2002-11-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: DHARAMSI, AKIL
APPLICANT: VEDADI, MASOUD
TITLE OF INVENTION: ESSENTIAL NOVEL BACTERIAL
FILE REFERENCE: ITT-330.01
CURRENT APPLICATION NUMBER: US/11/122,986
CURRENT FILING DATE: 2005-05-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/424,367
PRIOR FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,376
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PRIOR APPLICATION NUMBER: 60/423,758
PRIOR FILING DATE: 2002-11-05
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                                                                                                                                      22.9%;
Local Similarity 40.0%;
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APPLICATION NUMBER: 60/424,373
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    Mismatches

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Pred. No. 2.9e+0;
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Pred. No. 5.
                                                                                                                                                   Mismatches
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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  protein search, using
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1: /EMC Celerra SIDS3/ptodata/2/iaa/5 COMB.pep:*
2: /EMC Celerra SIDS3/ptodata/2/iaa/6 COMB.pep:*
3: /EMC_Celerra_SIDS3/ptodata/2/iaa/7 COMB.pep:*
4: /EMC_Celerra_SIDS3/ptodata/2/iaa/H COMB.pep:*
5: /EMC_Celerra_SIDS3/ptodata/2/iaa/PCTUS_COMB.pep:*
6: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*
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Maximum Match 100%
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US-09-343-623-8
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US-09-5510-10
US-08-390-509-10
US-08-390-790-10
US-08-390-790-10
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US-08-390-790-10
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US-08-151-574-3
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US-08-914-479A-11
                                                                                                                    Sequence 3, Application US/08151574 Patent No. 5436156
                                                                                                                                                                                                                                                                Query Match
Best Local
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                         GENERAL INFORMATION:
APPLICANT: Robert F.M. Van Gorcom
APPLICANT: Willem Van Harringsveldt
APPLICANT: Petrus A. Van Paridon
APPLICANT: Annemarie B. Veenstra
APPLICANT: Rudolf G.M. Luttin
APPLICANT: Gerardus Selten
  TITLE OF INVENTION:
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 Petrus A. Van Paridon
Annemarie E. Veenstra
Rudolf G.M. Luttin
Gerardus Selten
INVENTION: Cloning and EXIVENTION: Phytase
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| <b>4</b> 5       | 44              | 43              | 42              | 41              | 40                | 39                | 38               | 37               | 36               | 35               | 34               | S<br>S             | 32                | 31               | 30               | 29               | 28               | 27               |
|------------------|-----------------|-----------------|-----------------|-----------------|-------------------|-------------------|------------------|------------------|------------------|------------------|------------------|--------------------|-------------------|------------------|------------------|------------------|------------------|------------------|
| 26               | 26              | 26              | 26              | 26              | 26                | 26                | 26               | 26               | 26               | 26               | 26               | 27                 | 27                | 27               | 27               | 27               | 27               | 27               |
| 27.1             | 27.1            | 27.1            | 27.1            | 27.1            | 27.1              | 27.1              | 27.1             | 27.1             | 27.1             | 27.1             | 27.1             | 28.1               | 28.1              | 28.1             | 28.1             | 28.1             | 28.1             | 28.1             |
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| US-09-685-010-12 | US-09-233-510-9 | US-08-419-448-9 | US-08-196-940-3 | US-08-151-574-9 | US-09-978-309A-84 | US-09-978-309A-49 | US-09-977-827-14 | US-09-685-010-49 | US-09-343-623-13 | US-09-151-011-13 | US-08-486-839-13 | US-09-701-588C-107 | US-07-867-819D-75 | US-08-475-955-75 | US-09-977-827-15 | US-09-343-623-14 | US-09-151-011-14 | US-08-486-839-14 |
| Sequence         | Sequence        | Sequence        | Sequence        | Sequence        | Seguence          | Sequence          | Sequence         | Sequence         | Sequence         | Sequence         | Sequence         | Sequence           | Sequence          | Sequence         | Sequence         | Sequence         | Sequence         | sequence         |
| 12,              | ,               | 9               | ·               | 9               | 84,               | 49,               | 14,              | 49,              | 13,              | <u>ل</u> ا       | 13,              | 107                | 75,               | 75,              | 15,              | 14,              | 14,              | 14,              |
| Appl             | Appli           | Appli           | Appli           | Tiddy           | Appl              | Appl              | Appl             | Appl             | Appl             | Appl             | Appl             | , App              | Appl              | Appl             | Tddv             | Appl             | Appl             | Tddw             |

ALIGNMENTS

### FILE REFERENCE: 401112.404C2 CURRENT APPLICATION NUMBER: US/08/914,479A CURRENT FILING DATE: 1997-08-19 PRIOR APPLICATION NUMBER: 08/409,270 PRIOR FILING DATE: 1995-03-23 PRIOR APPLICATION NUMBER: 07/945,860 PRIOR FILING DATE: 1992-09-16 NUMBER OF SEO ID NOS: 19 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 11 LENGTH: 15 Sequence 11, Application US/08914479A Patent No. 6419932 GENERAL INFORMATION: APPLICANT: Dale, James B. TITLE OF INVENTION: ANTIGEN OF HYBRID M PROTEIN AND CARRIER TITLE OF INVENTION: FOR GROUP A STREPTOCOCCAL VACCINE OTHER INFORMATION: NH2-terminal fragment of M protein 1 OTHER INFORMATION: constructing antigens, which elicit OTHER INFORMATION: antibodies in an immunized animal TYPE: PRT ORGANISM: Artificial Sequence FEATURE: 2 DGQKVDEQFPQH 13 Similarity DARSVNGEFPRH Conservative 35.4%; 41.7%; 4. Score 34; DB Pred. No. 27; Mismatches 2 Length 15; Indels opsonic 0 Gaps

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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/151
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
24-MAY-1991
ATTORNEY/AGENT INFORMATION:
NAME: MUTABHIG, Kate H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 2461
TELECHOMMUNICATION INFORMATION:
TELEPHONE: 415-327-7250
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
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MOLECULE TYPE:
HYPOTHETICAL: N
FRAGMENT TYPE:
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APPLICANT:
APPLICANT:
ZIP: 20006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                         TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94025-3471
COMPUTER READABLE FORM:
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CITY: Washington
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Palane
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nes 6; Conserv
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STREET: 545 Middlefield Road, Suite 200
CITY: Menlo Park
                                                                                                                                                                                     ADDRESSEE:
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                                                                                                                                                                    2000 Pennsylvania Ave. N.W., Suite 5500
                                                                                                                   USA
                                                                                                                                                                                                                                        Annemarie E. Veenstra
Rudolf G.M. Luttin
Gerardus Selten
VENTION: Cloning and Expression of Microbial
VENTION: Phytase
                                                                                                                                                                                                                                                                                                                                 Robert F.M. Van Gorcom
Willem Van Hartingsveldt
Petrus A. Van Paridon
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                                                                                                                                                                                       Morrison & Foerster
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 66.7%;
Matches 6; Conservative
TELEFAX: 415-327-2951
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                    CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/688,57
APPLICATION NUMBER: 07/688,57
EILING DATE: 24 MAY-1991
ATTORNEY/AGENT INFORMATION:
NAME: MUTABhige, Kate H.
REGISTRATION NUMBER: 959
REFERENCE/DOCKET NUMBER: 2461
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDLIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0
                                                                                                                                                                                                                                                           SOFTWARE:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/233,510
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Rudolf G.M. Luttin
APPLICANT: Gerardus Selten
TITLE OF INVENTION: Cloning an
TITLE OF INVENTION: Phytase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Robert F.M. Van Gorcom
APPLICANT: Willem Van Hartingsveldt
APPLICANT: Petrus A. Van Paridon
APPLICANT: Annemarie E. Veenstra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide HYPOTHETICAL: NO FRAGMENT TYPE: N-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Murashige, Kate H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 10-APR-19
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 545 Middlefield Road, Suite 200
CITY: Menlo Park
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Murashige, Kate H. REGISTRATION NUMBER: 29,959
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
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                                                                                                                                                                                                        07/688,578
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                                                                                                               24615-20026.00
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RESULT 5
US-08-486-839-8
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                                              RESULT 6
US-09-151-011-8
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                                                                                                                                                                                                                                             US-08-486-839-8
                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/486
FILING DATE: 07 - June - 1995
ATTORNEY AGENT INFORMATION:
NAME: BATON, RODALD J.
REGISTRATION NUMBER: 29,281
REFERENCE/DOCKET NUMBER: 294-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 82-3550
TELEPAX: (516) 82-3582
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH 13 TELEPACE
           Sequence 8, Application US/09151011 Patent No. 6057142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8, Application US/08486839 Patent No. 5928928
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 33.3%;
Best Local Similarity 66.7%;
Matches 6; Conservative
                                                                                                                                                                          Query Match 31.2%;
Best Local Similarity 50.0%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V
SOFTWARE: DATA:
CURRENT APPLICATION DATA:
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                            TYPE: amino acids
STRANDEDNESS: unknown
TOPOLOGY: unknown
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TITLE OF INVENTION:
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HYPOTHETICAL:
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                                                                                                                                                                                                                                                          MOLECULE TYPE: protein HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: NO COUNTRY:
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: United States of America
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350 Jericho Turnpike
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Pred. No. 38;
1; Mismatches :
                                                                                                                                                                                             Score 30; DB Pred. No. 93;
                                                                                                                                                                            Mismatches
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US-09-343-623-8
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve
SOFTWARE:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/343,623
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/151,0:
APPLICATION NUMBER: US/09/151,0:
APPLICATION NUMBER: 10:
ATTORNEY/AGENT INFORMATION:
NAME: MOTTISS, ROBERT C.
REGISTRATION NUMBER: 294-32
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 822-3550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (516) 822-359
TELEFAX: (516) 822-3582
INFORMATION FOR SEQ ID NO: (
                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: FITTLE OF INVENTION: TITLE OF INVENTION: TUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 11791
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Hoffmann
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                     STATE: New Yo
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                                                                                                                                                    ZIP: 11758
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                                                                                                                                                                     COUNTRY:
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                                                                                                                                                                                                                                                                       A human chitinase, its recombinant production, its use for decomposing chitin, in therapy or prophylaxis against infection 16
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Pred. No. 93;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   294-32 DIV
                                                                       Version #1.30
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PRIOR APPLICATION DATA:

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RESULT 9
US-09-743-281C-136
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Best Local Similarity 5
""" ches 5; Conservat
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; LENGTH: 11
; TYPE: PRT
; ORGANISM: Autographa californica
US-09-977-827-8
            APPLICANT: Gaudernack, Gustav
APPLICANT: Eriksen, Jon Amund
APPLICANT: Moller, Mona
APPLICANT: Moller, Mona
APPLICANT: Seterdal, Ingvil
APPLICANT: Saeterdal, Ingvil
APPLICANT: Saeboe-Larsen, Stein
TITLE OF INVENTION: Antigenic Peptides Derived fro
FILE REFERENCE: 01702.401800
CURRENT APPLICATION NUMBER: US/09/743,281C
CURRENT FILING DATE: 2001-07-05
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ATTORNEY/AGENT INFORMATION:
NAME: BAZON, RONALD J.
REGISTRATION NUMBER: 29,281
REFERENCE/DOCKET NUMBER: 294-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 822-3580
TELEPAX: (516) 822-3582
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:

APPLICANT: Aerts, Johannes Maria F.G.

APPLICANT: Aerts, Johannes Maria F.G.

TITLE OF INVENTION: A human chitinase, its recombinant production, its use for decomp

TITLE OF INVENTION: chitin, its use in therapy or prophylaxis against infectious dis

FILE REFERENCE: Docket 294-32 DIVII/CON

CURRENT PELLICATION NUMBER: US/09/977,827

CURRENT FILING DATE: 2001-10-15

NUMBER OF SEQ ID NOS: 17

SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                           Sequence 136, Application US/09743281C Patent No. 7030211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 8, Application US/09977827 Patent No. 6896884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 31.2%;
Best Local Similarity 50.0%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 DGVDIDWEFP 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Mismatches
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Pred. No. 93;
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; NUMBER OF SEQ ID NOS: 231
; SOGTWARE: PatentIn version 3.0
; SEQ ID NO 136
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Homo sapiens, other or unknown US-09-743-281C-136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
                                                                                                                                      US-08-166-195A-23
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Verbion #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/166,195A
FILING DATE: 10 DEC 1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Sibley, Kenneth D.
NAME: Sibley, Kenneth D.
NAME: Sibley, Kenneth D.
                                                                  Matches
                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 5480799
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: O'Rand, Michael G.
APPLICANT: Widgren, Esther E.
APPLICANT: Richardson, Richard T.
APPLICANT: Lea, Isabel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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Best Local Similarity
Matches 5; Conserv
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: P.U. STREET: P.U. STREET: P.U. STREET: Charlotte
STATE: No. 5480799th Carolina
COUNTRY: USA
                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: peptide
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TITLE OF INVENTION: Sperm
TITLE OF INVENTION: Sperm
TITLE OF GROUENCES: 51
                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 919-881-3140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenneth D. Sibley
                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                     TYPE: amino acids
                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 5470/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23, Application US/08166195A
o. 5480799
1 GAKVDDRF 8
                                3 GOKVDEOF 10
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                                                                  Conservative
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                                                                                                                                                                                     single
                                                                                30.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sperm Antigen Corresponding to a Sperm Zona Binding Protein Autoantigenic Epitope
                                                                                Score 29; DB 1;
Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 29; DB 3;
Pred. No. 5e+05;
0; Mismatches
                                                                  Mismatches
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RESULT 11 US-08-436-772-23

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; TYPE: amino acid
STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-436-772-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 12
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                                                                                                                                                                                                                                                                                                                                                               Sequence 23, Application US/08436883B Patent No. 5820861 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 919-881-3140
TELEPAX: 919-881-3175
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                 APPLICANT: O'Rand, Michael G.
APPLICANT: Widgren, Bether E.
APPLICANT: Richardson, Richard T.
APPLICANT: Lea, Isabel
TITLE OF INVENTION: Sperm Antigen Corresponding to a Sperm
TITLE OF INVENTION: Cona Binding Protein Autoantigenic Epitope
NUMBER OF SEQUENCES: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Sibley, Kenneth D.
REGISTRATION NUMBER: 31,665
REFERENCE/DOCKET NUMBER: 54
TELECOMMUNICATION INFORMATION:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: O'Rand, Michael G.
APPLICANT: Widgren, Esther E.
APPLICANT: Richardson, Richard T.
APPLICANT: Lea, Isabel
TITLE OF INVENTION: Sperm Antigen Corresponding to a Sperm
TITLE OF INVENTION: Zona Binding Protein Autoantigenic Epitope
NUMBER OF SEQUENCES: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/436,772 FILING DATE: 08-MAY-1995 CLASSIFICATION: 435
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CITY: Charlotte
CTATE: No. 5814456th Carolina
                                                                  STREET: P.U. STREET: P.U. CITY: Charlotte
CITY: No. 5820861th Carolina
STATE: NGA
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Local Similarity 62.5%;
hes 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                     ADDRESSEE: Kenneth D. Sibley STREET: P.O. Box 34009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 GOKVDEOF 10
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P.O. Box 34009
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Pred. No. 1.2e+02;
2; Mismatches 1;
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LENGTH: 10 amino acids
TYPE: amino acid
TYPE: samino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-436-883B-23
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; MOLECULE TYPE:
; HYPOTHETICAL:
US-08-486-839-12
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Best Local Similarity 62.5%;
Matches 5; Conservative
                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/486
FILING DATE: 07 - June - 1995
ATTORNEY/AGENT INFORMATION:
NAME: Baron, Ronald J.
REGISTRATION NUMBER: 29,281
REFERENCE/DOCKET NUMBER: 294-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 822-3550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 919-881-3175
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                          INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: A hutile of INVENTION: procing in the INVENTION: in the INVENTION of SEQUENCES: 16 CORRESPONDENCE ADDRESS:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/436,883B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: 919-420-2200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 350 Je
CITY: Jericho
STATE: New Yor
                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy diak
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Sibley, Kenneth D.
REGISTRATION NUMBER: 31,665
REFERENCE/DOCKET NUMBER: 5470-73C
                                                                                                                                                                                                                                                                                                                                          SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM:
                                                    TOPOLOGY:
                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Sibley,
                                                                                                       ENGTH:
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5928928
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                                                                                    amino acid
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                                                                                                           11 amino acids
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                                                                                                                                                             (516) 822-3582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      United States of America
                                                    unknown
             .. protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-MAY-1995
                                                                       unknown
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                                                                                                                                                                                                                                                                                                         US/08/486,839
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Pred. No. 1.2e+02
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US-09-151-011-12
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                                                                                                                                                                                        RESULT 15
                                                                                                                                                                                                                                                                                                                                                                                                    US-09-151-011-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: MORTISS, Robert C.
REGISTRATION NUMBER: 42,910
REFERENCE/DOCKET NUMBER: 294-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 822-3550
TELEPAX: (516) 822-3582
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 30.2%;
Best Local Similarity 50.0%;
Matches 5; Conservative
                                                                                       Sequence 12, Application US/09343623
Patent No. 6303118
GENERAL INFORMATION:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 12, Application US/09151011 Patent No. 6057142
                                                                                                                                                                                                                                                                                                                    Query Match 30.2%;
Best Local Similarity 50.0%;
Matches 5; Conservative
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GENERAL INFORMATION:
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TITLE OF INVENTION: There
TITLE OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hoffmann &
               TITLE OF INVENTION: 1
TITLE OF INVENTION: 1
TITLE OF INVENTION: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/151,011
FILING DATE: 10 - September - 1998
ATTORNEY/AGENT INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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STREET: SYOBBET
CITY: SYOBBET
TATE: New York
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CORRESPONDENCE ADDRESS:
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             A human chitinase, its recombinant production, its use for decomposing chitin, its use in therapy or prophylaxis against infection diseases 16
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Production, Its Use For Decomposing Chitin, Its Use in
Therapy or Prophylaxis Against Infection Diseases.
16
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Pred. No. 1.4e+02.
2; Mismatches
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Pred. No. 1.4e+02;
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Search completed: May 30, 2006, 15:09:24 Job time: 19.3333 secs
                                                                                         Query Match
Best Local Similarity
Watches 5; Conserve
                                                                                                                                                        US-09-343-623-12
                                                                                                                                                                                                                                                              TELEPHONE: (516) 822-35
TELEFAX: (516) 822-3582
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 11758
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                    MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/343,623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Hof:
STREET: 350 Je:
CITY: Jericho
STATE: New Yorl
                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.30
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                                                     DGLNLDWQYP 11
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350 Jericho Turnpike
                                                                                                        Conservative
                                                                                                                                                                                                unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        United States of America
                                                                                                                                                                    : protein
NO
                                                                                                                   30.2%;
                                                                                                        Mismatches
                                                                                                                   Score 29; DB 2;
Pred. No. 1.4e+02;
                                                                                                                               Length 11;
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Result
No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Searched:
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Maximum Match
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                  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed,
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                  protein search, using sw
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length: 18
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Match
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1: pir1:*
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A61577
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A28144
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S71948
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alpha-macroglobuli
protein P2 - curle
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| <b>6.</b>  |
|  |

ALIGNMENTS

#### protein Pi1 - golden needle mushroom (fragment) C;Species: Flammulina velutipes (golden needle mushroom) C;Date: 12-Nov-1999 #sequence\_revision 12-Nov-1999 #text\_change 09-Ju1-2004 C;Date: 12-Nov-1999 #sequence\_revision 12-Nov-1999 #text\_change 09-Ju1-2004 C;Accession: A59137 R;Sakamoto, Y.; Ando, A.; Tamai, Y.; Miura, K. submitted to the Protein Sequence Database, November 1999 A;Bescription: Differences of proteins expressed in the fruiting dikaryon ar A;Reference number: A59137 A;Accession: A59137 A;Cross-references: UNIPARC:UPI0000176DF0 A;Experimental source: patient SS, IL-2R+ synovial T-cells A;Note: sequence extracted from NCBI backbone (NCBIP:113270) C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: T-cell receptor T-cell receptor beta chain V region (CDR3 junction, clone Vbeta20.1) - human (fragment) C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 21-Jan-1994 #sequence\_revision 18-Nov-1994 #text\_change 30-May-1997 C;Accession: H49048 R;Sioud, M.; Kjeldsen-Kragh, J.; Suleyman, S.; Vinje, O.; Natvig, J.B.; Forre, O. Bur. J. Immunol. 22, 2413-2418, 1992 A;Title: Limited heterogeneity of T cell receptor variable region gene usage in juvenile A;Reference number: A49048 MUID:92387250; PMID:1387614 A;Accession: H49048 A;Status: preliminary; not compared with conceptual translation A;Status: preliminary A;Molecule type: protein A;Residues: 1-18 <3AK> A;Cross-references: UNIPROT:Q7M4W6; UNIPARC:UPI000017CB27 밁 Ş 문 á A59137 RESULT 2 A; Molecule type: mRNA A; Residues: 1-17 <SIO> H49048 Matches Query Match Matches Query Match Best Local Local Similarity 11 PYTAP 15 N æ σ Similarity 5; Conservat PYTSP ATNIFPYT 13 AWTEFPYT 15 Conservative Conservative 27.8**%**; 80.0**%**; 27.8**%**; 62.5**%**; 0 Score 27; DB 2; Le Pred. No. 3.8e+02; 1; Mismatches 0; Score 27; DB 2; Pred. No. 3.5e+02; Mismatches in the fruiting dikaryon and the non-fi Length 18; Length 17; Indels Indels 0 0 Gaps Gaps 0 0

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R;Binette, J.P.; Binette, M.B.
submitted to the Protein Sequence Database, October 1996
A;Description: The proteins of kidney and gallbladder stones
A;Reference number: A58501
A;Accession: B58502
                                                                                                                                                                                                                                                                                      C;Date: 07-A;
C;Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                36K kidney stone protein - unidentified bacterium (fragment)
C;Species: unidentified bacterium
C;Date: 07-Feb-1997 #sequence_revision 07-Feb-1997 #text_change 10-Jul-1998
C;Accession: B58502
R;Binette, J.P.; Binette, M.B.
                                                                                                                                                     A;Reference number: S18530; MUID:92079886; PMID:1840640
A;Accession: S18534
                                                                                                                                                                                                              Mol. Gen. Genet. 230, 120-128, 1991
A;Title: Cloning and sequence analysis of genes involved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPARC:UPI000017ABD0
A;Experimental source: human kidney stone containing
A;Note: tentative identification of 8-Tyr and 9-Thr
                                           A;Cross-references: UNIPARC:UPI000017AD81; A;Note: the authors translated the codon CT
                                                                                       A; Molecule type: DNA
A; Residues: 1-17 <HAY>
                                                                                                                                   A;Status: preliminary
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A; Residues: 1-18 <SIN>
A; Cross-references: UNIPROT: Q7M198; UNIPARC: UPI000017AE13
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R;Sinha, U.; Wolz, S.A.; Lad, P.J.
Int. J. Biochem. 23, 979-984, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24k serine proteinase (EC 3.4.21.-) - Streptomyces fradiae (fragment) C;Species: Streptomyces fradiae (C;Date: 28-Oct-1994 #sequence_revision 28-Oct-1994 #text_change 09-Jul-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C; Keywords: extracellular protein; hydrolase; serine
                                                                                                                                                                                                 methyltransferases.
                                                                                                                                                                                                                                          Haydock, S.F.; Dowson, J.A.; Dhillon, N.; Roberts, G.A.; Cortes, J.; Leadlay, P.F.
                                                                                                                                                                                                                                                                                                 ;Species: Saccharopolyspora erythraea
;Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text_change 07-Apr-1994
Query Match
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Similarity 41.7%;
5; Conservation
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  24.78;
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1; Mismatches
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Pred. No. 5.6e+02;
  24;
                                         L; EMBL:X60379
CTG for residue 12 as
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R; Leterme, S.; Boutry, M.
Plant Physiol. 102, 435-443, 1993
A;Title: Purification and preliminary characterization of mitochondrial complex I (NADH: A;Reference number: PQ0775; MUID:94151437; PMID:8108509
A;Accession: PQ0786
                                                               RESULT 8
S20322
gluten - wheat
C;Species: Triticum
C;Date: 19-Mar-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Wiederrecht, G.; Hung, S.; Chan, H.K.; Marcy, A.; Martin, M.; Calaycay, J.; J. Biol. Chem. 267, 21753-21760, 1992
A;Title: Characterization of high molecular weight FK-506 binding activities A;Reference number: A45115; MUID:93016131; PMID:1383226
A;Accession: B45115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  peptidylprolyl isomerase (EC 5.2.1.8) FKBP51 - human (fragment) N;Alternate names: FK506-binding protein FKBP51; peptidylprolyl C;Species: Homo sapiens (man) C;Date: 30-Apr.1993 #sequence_revision 18-Nov-1994 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:07M2G3; UNIPARC:UPI000017CDA8
C;Comment: Complex I, mitochondrial NADH-ubiquiquinone reductase,
ranging from 5K to 75K.
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A;Experimental source: JURKAT cells
A;Note: sequence extracted from NCBI backbone (NCBIP:116748)
C;Keywords: cis-trans-isomerase; cyclosporin A binding
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C; Genetics:
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Best Local
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Pred. No. 1.2e+03;
Pred. --hem 2;
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Pred. No. 1.5e+03;
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0; Mismatches 3;
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aestivum (common wheat)
#sequence\_revision 19-Mar-1997 #text\_change 09-Jul-2004

C;Accession: S20322 R;Fukudome, S:; Yoshikawa, M. FEBS Lett. 296, 107-111, 1992 A;Title: Opioid peptides derived from wheat gluten: their isolation and characterization A;Reference number: S20322; MUID:92111773; PMID:1309704

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R;Chen, H.C.; Wintz, H.; Weil, J.H.; Pillay, D.T.N.
Nucleic Acids Res. 16, 10372, 1988
A;Title: Nucleotide sequence of chloroplast CF1-ATPase epsilon-subunit and A;Reference number: S01903; MUID:89057486; PMID:2904134
A;Accession: S01904
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PN0170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Accession: PN0170
R;Fukaya, N.; Chow, L.P.; Sugiura, Y.; Tsugita, A.; Ueno, Y.; Tabuchi, K.
submitted to JIPID, May 1994
A;Description: Two dimensional polyacrylamide gel electrophoresis of Fusarium
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S01904
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A; Accession: PN0170
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C;Date: 05-Aug_1994 #sequence_revision 06-Jan-1995 #text_change 09-Jul-2004
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A;Genome: ch
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A; Residues: 1-13 < CHE >
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66.7%;
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Pred. No. 1.9e+03;
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Pred. No. 1.8e+03;
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RESULT 12
PA0087
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A;Description: Two dimensional polyacry A;Reference number: PA0051
A;Accession: PA0099
                                                                                                                                                                                                                                                                                phenotipic variation protein - fungus (Fusarium sporotrichioides) (fragment) (;Species: Fusarium sporotrichioides C;Species: Fusarium sporotrichioides C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004 C;Accession: PA0099 R;Chow, L.P.; Fukaya, N.; Sugiura, Y.; Ueno, Y.; Tabuchi, K.; Tsugita, A.
                                                                                                                                                                                                                                                                                                                                                                               PA0099
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A;Description: Two dimensional polyacrylamide gel electropheresis of Fusarium sporotrich A;Reference number: PA0051
A;Accession: PA0056
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Chow, L.P.; Fukaya, N.; Sugiura, Y.; submitted to JIPID, October 1994
A;Description: Two dimensional polyacry
A;Reference number: PA0051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cytochrome c2 - fungus (Fusarium sporotrichioides) (fragment)
C;Species: Fusarium sporotrichioides
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_char
C;Accession: PA0087
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C;Species: Pusarium sporotrichioides
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 23-Mar-2001
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A;Residues: 1-15 <CHO>
A;Cross-references: UNIPROT:07M4Z7; UNIPARC:UPI000017B3F8
C;Keywords: electron transfer; heme; photosynthesis
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A;Residues: 1-15 <CHO>
A;Cross-references: UNIPARC:UPI000017B408
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C;Accession: C84063
R;Takami, H.; Nakasone, K.; Takaki, Y.; Maeno, G.; Sasaki, R.; Masui, N.; Fuji, F.; Hira Nucleic Acids Res. 28, 4317-4331, 2000
A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodurans and A;Reference number: A83650; MUID:20512582; PMID:11058132
A;Accession: C84063
A;Status: preliminary
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-17 <STO>
A;Cross-references: UNIPROT:Q9K7Q4; UNIPARC:UPI00000C4168; GB:AP001518; GB:BA000004; NID
A;Experimental source: strain C-125
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c;Species: Bos prinigenius taurus (cattle)
C;Date: 28-Aug-1989 #sequence_revision 28-Aug-1989 #text_change 09-Jul-2004
C;Accession: A2814
R;Redman, K.L.; Rechsteiner, M.
J. Biol. Chem. 263, 4926-4931, 1988
A;Title: Extended reading frame of a ubiquitin gene encodes a stable, conserved, basic parameters are protein approached to the protein A;Accession: A28144; MUID:88169619; PMID:2832412
A;Accession: A28144; MUID:88169619; PMID:2832412
A;Accession: A28144
A;Molecule type: protein
A;Cross-references: UNIPROT:097577; UNIPARC:UPI0000177161
C;Superfamily: ubiquitin / rat ribosomal protein S27a; ribosomal protein biosynthesis
Search completed: May 30, 2006, 15:08:21 Job time : 12 secs
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## SUMMARIES

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7 Q1-MAY-2000, integrated into UniProtKB/TrEMBL.
7 Q1-MAY-2000, sequence version 1.
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PubMed=15019625; DOI=10.1016/j.ympev.2003.08.002;
Paton A., Springate D.A., Sudde S., Otieno D., Grayer R., Harley |
Paton A., Springate D.A., Sudde S., Otieno D., Grayer R., Harley |
Willis F., Simonds M.S.J., Powell M.P., Savolainen V.;
"Phylogeny and evolution of basils and allies (Ocimeae, Labiatae)
based on three plastid DNA regions.",
based on three plastid DNA regions."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=7375;
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prina (Greenbottle fly) (Australian sheep blowfly).
Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
andopterygota; Diptera; Brachycera; Muscomorpha; Oestroidea;
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                                                                                                     PRELIMINARY;
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58.3%;
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; Mismatches
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Pred. No. 2
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                                                                                                        PRT;
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ches 2;
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Attribution-NoDerivs License
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RESULT
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Q7M4W6_FLAVE PRELIMINARY;
Q7M4W6;
15-DEC-2003, integrated into
15-DEC-2003, sequence versio
07-FEB-2006, entry version 6
Protein Pil (Fragment).
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SEQUENCE 18 AA; 199
               07 NEUCR
Q75007 NEUCR PRELIMINARY; PRT; 15 AA.
Q75007;
15-DEC-2003, integrated into UniProtKB/TrEMBL
15-DEC-2003, sequence version 1.
07-FEB-2006, entry version 8.
Predicted protein.
ORFNames=NCU09752.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Flammulina velutipes.
Eukaryota; Fungi; Basidiomycota; Hymenomycetes; Homobasidiomycetes; Agaricales; Tricholomataceae; Flammulina.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
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Nucleic Acids Res. 30:20
Neurospora
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sakamoto Y., Ando A., Tamai Y., Miura K.;
Submitted (NOV-1999) to the PIR data bank
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STRAIN=UTEX 485;
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Eukaryota; Fungi; I Sordariomycetidae;

Ascomycota; Pezizomycotina; Sordariomycetes; Sordariales; Sordariaceae; Neurospora.

NCBI\_TaxID=5141;

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SOTTER COCCEPT AND SOTTER COCCEP
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RA Jaffe D., FitzHugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B.,
RA Elkins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M.,
Qui D., Ianakiev P., Bell-Pedersen D., Nelson M.A.,
RA Qui D., Ianakiev P., Bell-Pedersen D., Nelson M.A.,
RA Zelter A., Schulte U., Kothe G.O., Jedd G., Mewes H.-W., Staben C.,
RA Marcotte E., Greenberg D., Roy A., Foley K., Naylor J.,
RA Marcotte E., Bielke C., Rudd S., Foley K., Naylor J.,
RA Mauceli E., Bielke C., Rudd S., Frishman D., Krystofova S.,
RA Macino G., Catcheside D.E.A., Li W., Pratt R.J., Osmani S.A.,
RA Agamusen C., Glass N.L., Orbach M.J., Berglund J.A., Voelker R.,
RA Paulsen C., Plamann M., Seiler S., Dunlap J.C., Radford A., Aramayo R.,
RA Natvig D.O., Alex L.A., Mannhaupt G., Ebbole D.J., Freitag M.,
RA Paulsen I., Sachs M.S., Lander E.S., Nusbaum C., Birren B.W.;
"The genome sequence of the filamentous fungus Neurospora crassa.";
RI Nature 422:859-868(2003).
Query Match
Best Local S
Matches 5
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Matches 4
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"Two new extracellular serine proteases
Int. J. Biochem. 23:979-984(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24k serine proteinase (EC 3.4.21.-) (Fragment). Streptomyces fradiae.
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15-DEC-2003, sequence version 1.
07-FEB-2006, entry version 7.
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Q7M198;
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                                                                                                                                                                                                                                     Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=92155439; PubMed=1786859; DOI=10.1016/0020-711X(91)90133-8;
                                                                                                                                                                                                                                                                                                                                                                                                                       PROTEIN SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Streptomycineae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Distributed under the Creative Commons Attribution-NoDerivs License
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STRAIN=74-OR23-1A / FGSC 987;
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1832 MW; 012D0180A8C7089D
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2; Mismatches
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Best Local Similarity
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Q9UR63;
                                                                                                              MEDLINE=96409246; PubMed=8814228; DOI=10.1016/0167-4838(96)00073-8; Chen J.S., Saxton J., Hemming F.W., Peberdy J.F.; "Purification and partial characterization of the high and low molecular weight form (S- and F-form) of invertase secreted by Aspergillus nidulans."; Biochim. Biophys. Acta 1296:207-218(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAY-2000, integrated into UniProtKB/TrEMBL.
01-MAY-2000, sequence version 1.
07-FEB-2006, entry version 11.
Beta-D-FRUCTOFURANOSIDE FRUCTOHYDROLASS 60 kDa
                                            Copyrighted by the UniProt Consortium, see http://www.uniprot.org/termsDistributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Differential extraction and protein sequencing differences in patterns of primary cell wall pro J. Biol. Chem. 272:15841-15848(1997).
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25-OCT-2005, sequence versi
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40 kDa cell wall protein (F
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MEDLINE=96409246;
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Eurotiales; Trichocomaceae; Emericella.
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1 >15 40 kDa cell wall protein.
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PubMed=9188482; DOI=10.1074/jbc.
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C., Bolwell G.
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RESULT 11
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27-SEP-2005, integrated into Unit
27-SEP-2005, sequence version 1.
07-FEB-2006, entry version 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Q3ZEV4_9CHAR
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                                                                                                                                                           STRAIN=CCMP1319;
PubMed=14681369; DOI=10.1093/nar/gkh102;
Gueneau de Novoa P., Williams K.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mitochondrion.
NON_TER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATP synthase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Copyrighted by the UniProt Consortium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Condor 107:514-526(2005).
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NCBI_TaxID=52970;
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                                                                                                                           "The tmRNA website: reductive
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(Fragment)
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Pred. No. 8.2e+03
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07-FEB-2006,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.
Soura V., Semikhodskii A.G.,
Submitted (JUN-2003) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Phodopus sungorus (Striped hairy-footed hamster) (Djungarian hamster). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Muroidea; Cricetidae; Cricetinae; Phodopus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQÜENCE
                            Pereira S.L., Baker A.J.;
"Multiple gene evidence fi
ancestral morphological si
Scolopacidae).";
                                                                                                                               Eukaryota; Metazoa; Chordata; (Archosauria; Aves; Neognathae;
                                                                                                                                                              Mitochondrion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NON
                                                                                                                                                                            Tringa flavipes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=10044;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Per2 protein
                  Condor 107:514-526(2005)
                                                                                         NUCLEOTIDE SEQUENCE.
                                                                                                                   NCBI_TaxID=161739;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AF550355; AAQ12671.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AY316536; AAP87362.1;
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                                                                                                                                                                                                                                      sequence version 1.
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                                                                                                                                  Chordata; Craniata; Vertebrata; Euteleosto
Neognathae; Charadriiformes; Scolopacidae;
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Pred. No.
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STRAIN=C57BL/6J;

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RESULT 15
Q3TW38 MOUSE
ID Q3TW38;
DT 11-OCT-2005, i
DT 11-OCT-2006, e
DE OSteoclast-lik
DE clone:1420034D
DE interacting pr
GN Name=Rbms1;
OS Mus musculus
OC Eukaryota; Met
OC Murroidea; Muri
OX NCBI_TaxID=100
RN [1]
RP NUCLEOTIDE SEQ
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NON TER 1
SEQUENCE 16 F
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Q65YZ6_HORSE
Q65YZ6;
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Distributed
                                                                                                                                                                                               Osteoclast-like cell cDNA, RIKEN full-length enriched library, clone:I420034D23 product:RNA binding motif, single stranded interacting protein 1, full insert sequence. (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE.
Hasegawa T., Sato F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Crani
Mammalia; Eutheria; Laurasiatheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-OCT-2004, sequence version 07-FEB-2006, entry version 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>ვ</u>
                                                                                                                                                                                                                                                                        11-OCT-2005, sequence version 1.
07-FEB-2006, entry version 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Equus caballus (Horse).
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     NUCLEOTIDE SEQUENCE
                               NCBI_TaxID=10090;
[1]
                                                                                                                                                   Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SLFPLLFTAP
                                                                     ; Metazoa; Chordata; Craniata;
Eutheria; Euarchontoglires; Gl
Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ping of equine chromosomes corresponding to HSA (MAR-2004) to the EMBL/GenBank/DDBJ databases.
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17
17 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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50.0%;
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Pred. No. 1.2e
1; Mismatches
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i, Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                             PRT;
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heria; Perissodactyla;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hirota K.;
                                                                                                  Glires; Rodentia; Sciurognathi;
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1.2e+04;
                                                                                                                              Vertebrata; Euteleostomi;
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RX DEMMENSIALINOUS MELLONIS CONTROL OF THE PROBLEM OF THE PROCESS OF THE PROBLEM 
     Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S., Okazaki G., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H. Yagi K., Tomaru Y., Hasegawa Y., Nogami A., Schonbach C., Gojobori Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J., Schriml L.M., Kanapin A., Matsuda H., Batalov S., Beisel K.W., Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins Dalla E., Dragani T.A., Fletcher C.F., Forrest A., Frazer K.S., Gassterland T., Gariboldi M., Gissi C., Godzik A., Gough J., Gassterland T., Gariboldi M., Gissi C., Godzik A., Gough J., Gassterland S., Gusthncich S., Hirokawa N., Jackson I.J., Jarvis E.D., Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L., Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A., Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RIKEN Genome Exploration Research Group, and Genome Science (Genome Network Core Team) and the FANTOM Consortium; "Antisense Transcription in the Mammalian Transcriptome.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                         "Antisense Transcription in Science 309:1564-1566(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PubMed=16141073; DOI=10.1126/science.1112009;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=C57BL/6J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "The transcriptional landscape of the mammalian genome."; Science 309:1559-1563(2005).
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RA RAVABÍT., Reed J.C., Reed D.J., Reid J., Ring B.Z., Ringwald M., RA Sandelin A., Schneider C., Semple C.A., Serou M., Shimada K., RA Sultana R., Takenaka Y., Taylor M.S., Teasdale R.D., Tomita M., RA Verardo R., Wagner L., Wahlestedt C., Wang Y., Watanabe Y., Wells C., RA Vilming L.G., Wynshaw-Boris A., Yanagisawa M., Yang I., Yang Y., Zavolan M., Zhu Y., Zimmer A., Carninci F., Hayatsu N., RA Hirozane-Kishikawa T., Konno H., Nakamura M., Sakazume N., Sato K., RA Shiraki T., Waki K., Kawai J., Aizawa K., Arakawa T., Fukuda S., RA Hara A., Hashizume W., Imoteni K., Ishii Y., Itoh M., Kagawa I., RA Yasunishi A., Yoshino M., Waterston R., Lander E.S., Rogers J., A Birney E., Hayashizaki Y., Tanalysis of the moving """
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RX KAWAI J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
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RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Sasaki H., Sato K., Wang K.H., Weitz C., Whittaker C., Wilming L.,
RA Mynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
Hayashizaki Y.,
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Kawai J., Kojima ..., Ohno M., U
Nishiyori H., Nomura K., Ohno M., U
Shibata K., Shiraki T., Tagami M.,
Muramatsu M., Hayashizaki Y.,
Submitted (MAR-2004) to the EMBL/Ge
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Genome R
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Nature 420:563-573(2002).
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                                                                                        GO:0005634; C:nucleus; RCA.
GO:0003690; F:double-stranded DNA binding; RCA.
GO:0003723; F:RNA binding; RCA.
GO:0003697; F:single-stranded DNA binding; RCA.
GO:0003697; F:DNA replication; RCA.
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Adj38596 HSV-4 Gly
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# ALIGNMENTS

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RESULT 1
ADR10610
Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody;
                                                                                                                                                                                                                                                      Cat IgE epitope recognised by monoclonal antibody 3.76, SEQ ID
                                                                                                                                                          16-JAN-2003; 2003US-0440472P.
                                                                                                                                                                                    05-AUG-2004.
                                                                                                                                                                                                               Felis catus
                                                                                                                                                                                                                                                                      21-OCT-2004
                                                                                                                                                                                                                                                                                   ADR10610;
                                                                                                                                                                                                                                                                                                ADR10610 standard; peptide;
                                                                                                                                            (UYNC-) UNIV NORTH CAROLINA STATE
                                                                                                                                                                      15-JAN-2004; 2004WO-US003566
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                                                                                                                                                                                                                                                                     (first entry)
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Hammerberg B;

WPI; 2004-593545/57.

Novel antibody that specifically binds to for testing an allergen reactivity of IgE or treating asthma or anaphylactic shock. mammalian IgE epitope, useful sample, detecting mammalian I IgE

Example 6; Page 9; 14pp; English.

The present invention relates to a novel monoclonal antibody (I) that specifically binds to a mammalian IGE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IGE, e.g. dog IGE. (I) is useful for testing an allergen reactivity of an IGE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy peanut and corn allergens. The sample is a biological sample collected from a dog, cat or horse. (I) is also useful for detecting mammalian IGE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal antibodies recognise epitopes on canine IGE corresponding to amino acid residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the

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RESULT 2
ADRILGED9
ID ADRILGED9
ID ADRILGED9
ID ADRILGED9
XX ADRIL
                                                            The present invention relates to a novel monoclonal antibody (I) that CC specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. CC (I) is useful for testing an allergen reactivity of an IgE sample. The callergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut CC and corn allergens. The sample is a biological sample collected from a CC dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal CC antibodies recognise epitopes on canine IgE corresponding to amino acid CC enihorse, pig and human by 5.91 and 3.76 monoclonal cc canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from CC cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and 3.76 were observed to have good cross-reactivity with the epsilon-chain CC ceither pig or human epsilon-chains of IgE from cat and horse, but did not exhibit cross-reactivity with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian Igor treating asthma or anaphylactic shock.
   Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hammerberg B;
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Pred. No. 5.7e-09;
; Mismatches 0;
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Best Local Similarity

89.7%; 100.0%;

Score 87; Pred. No.

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Length 18

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Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sheep IgE epitope recognised
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                                                                                                              Score 55; DB 8;
Pred. No. 0.067;
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                                                                                                                                 Length 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to a novel protein, human protein 36-41, and the DNA sequence encoding it. The protein of the invention may be useful for the treatment of diseases such as arrhythmia, asthma and dementia. The present sequence is the amino acid sequence of a peptide fragment of human protein 36-41 which was used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New human macroprotein-36.41, encoding polynucleotide, antagonist recombinant production, useful for treating dementia, arrhythmia, and digestive ulcers.
                                                                                                                                     Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody;
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                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15 AA;
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16-JAN-2003; 2003US-0440472P
                                                                                                                                                                         Horse IgE epitope recognised by monoclonal antibody 3.76,
                                                                                                                                                                                                                              ADR10611;
                                                                                                                                                                                                                                                      ADR10611 standard; peptide; 18
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                         15-JAN-2004; 2004WO-US003566.
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                                                                                                  Equus caballus
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                                                                                                                                                                                                                                                                                                                   MDGKMQPNSFPWQSP 15
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40.0%;
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Pred. No. 59;
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                                                                                                                                                                                                              Gershwin LJ,
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                                                                                                                                                                                                              Pettigrew HD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 35; DB 8; Le
Pred. No. 1.6e+02;
1; Mismatches 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       heavy chain immunogenic peptide P5
                                                                                                                                                                                                                 Kalina WV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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Immunogenic composition comprising an isolated equine immunoglobulin polypeptide that induces production of antibodies which specifically to equine immunoglobulin E.

E bind

Example 1; Page 8; 14pp; English.

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CC antibody that specifically binds to equine immunoglobulin (1g)E), the six CC polypeptides are not explicitly identified in the specification. Also CC included are a composition comprising an antibody that specifically binds to equine IgE made by the process of immunising an animal with a polypeptide at least 80% identical to (S1)-(S6), an antibody that specifically binds to equine IgE made by the process of immunising an CC antibody that specifically binds to equine IgE (involving immunising an CC amino acid sequence of the polypeptide is at least 80% identical to (S1)-(S6), making an CC amino acid sequence of the polypeptide is at least 80% identical to (S1)-(CC (S6)), and collecting antiserum from the antibody and means CC of equine IgE in a biological sample comprising the antibody and means CC for detecting specific binding of the antibody to equine IgE. The CC sample (serum) which involves contacting the sample with the antibody. CC thus forming an antigen/antibody complex, and detecting the presence or complex can be detected. The complex is detected using a second labelled complex can be detected. The complex is detected using a second labelled complex can be detected. The complex is detected using a second labelled complex is a Horse immunoglobulin E, IgE, heavy chain immunogenic cc peptide from the middle portion of the C2 region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches
                                                                                                                          19-JUL-2002; 2002US-0397265P.
19-JUL-2002; 2002US-0397477P.
19-JUL-2002; 2002US-0397479P.
03-MAR-2003; 2003US-0451903P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human herpesvirus Synthetic.
   Morham
                                                                                                                                                                                                                                                                                         21-JUL-2003; 2003WO-US022828.
                                                                                                                                                                                                                                                                                                                                                             29-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2004009027-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Virucide; HSV infection; antiviral; late domain motif; mutein; mutant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HSV-4 Glycoprotein B late domain motif mutant peptide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an immunogenic composition comprising an isolated polypeptide having an amino acid sequence that is at least identical to 6 (S1-S6), 15 amino acid peptide sequences derived from equine immunoglobulin E (the composition induces production of an
                                                                (MYRI-) MYRIAD GENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADJ38597 standard; peptide; 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 VDGQKATNIFP 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IDGQKVDEQFP 12
Zavitz K,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "Wild-type residue replaced with Ala"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             35.1%;
   Hobden A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 34; DB 7;
Pred. No. 1.9e+02;
1; Mismatches 4
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Example 13; Page 66; 74pp; English

Use of cells displaying herpes simplex virus (HSV) altered phenotype for the manufacture of a medicament for treating

hsv infection

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RESULT 8
ADJ38596
ID ADJ3
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           contain a nucleic acid encoding a mutant HSV protein or a nucleic acid encoding the polypeptide sufficient for virus-like particle assembly but devoid of late-domain motifs. The nucleic acid is within an HSV genome. The genome is devoid of late domain motifs capable of effecting viral budding. The composition also comprises a compound capable of interfering with the protein-protein interaction between a host cell protein capable of binding a late domain motif and a HSV protein containing a late domain motif. The composition further comprises another HSV protein or its immunogenic fragment, and/or a nucleic acid encoding the other HSV protein or the immunogenic fragment. The present sequence is a mutant HSV peptide, derived from the wild-type peptide ADJ38596, used to illustrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to cells displaying herpes simplex virus (HSV) altered budding phenotype which are useful for the manufacture of a medicament for treating HSV infection. The medicament further comprises an adjuvant capable of stimulating cellular immune response e.g. IL-2, IL-4, IL-12, IL-18 or gamma-interferon. The cells are human cells and
                                                                                                                                                                                       19-JUL-2002; 2002US-0397265P.
19-JUL-2002; 2002US-0397479P.
19-JUL-2002; 2002US-039749P.
03-MAR-2003; 2003US-0451903P.
                                                                                                                        Morham S,
                                                                                                                                                                                                                                                                    21-JUL-2003; 2003WO-US022828
                                                                                                                                                                                                                                                                                                    29-JAN-2004.
                                                                                                                                                                                                                                                                                                                                 WO2004009027-A2
                                                                                                                                                                                                                                                                                                                                                             Human herpesvirus
                                                                                                                                                                                                                                                                                                                                                                                              Virucide; HSV infection; antiviral; late domain motif
                                                                                                                                                                                                                                                                                                                                                                                                                            HSV-4 Glycoprotein B late domain motif wild-type peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADJ38596;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADJ38596 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 13; Page 66; 74pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Use of cells displaying herpes simplex virus (HSV) altered phenotype for the manufacture of a medicament for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-123282/12.
                                                                                                                                                       (MYRI-) MYRIAD GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6 ATNIFPYTAPGK 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTV-PATAPGK
                                                                                                                        Zavitz K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  34.5%;
66.7%;
                                                                                                                          Hobden A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 33.5; D
Pred. No. 2.3e
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 15;
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HSV infection
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RESULT 9
ADW78189
ID 78189
ID 78189
ID 78189
ID 78189
AC ADW7

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vasotropic; hypotensive; antiarteriosclerotic; antiinflammatory; antiallergic; immunosuppressive; antibacterial; antiasthmatic; dermatological; antidabetic; gastrointestinal-gen.; antiulcer; thrombolytic; neuroprotective; ophthalmological; antirheumatic; antipyretic; uropathic; antipsoriatic; hepactoropic; antianemic; muscular-gen.; thyromimetic; antithyroid; gynecological; nephrotropic; hepatotropic; virucide; anti-HIV; anabolic; hypertensive; anticonvulsant; endocrine-gen.; neuroleptic; nootropic; antiparkinsonian; anticonvulsant; antidepressant; antiaddictive; seedative; hypnotic; CNS-gen.; antidepressant; antiaddictive; seedative; hypnotic; CNS-gen.; anticonvulsant; antiaddictive; seedative; hypnotic; cipsonian; anticonvulsant; metabolic disorder; neuropathy; sleep disorder.
                                                                                Identification of compound for treating e.g. cancer by culturing cells expressing target protein in the presence of growth factor, cytokine, tumor promoter or oncogene and assessing binding after contacting with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human metabolic therapy target peptide PAGT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-APR-2005
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                                                 the compound
                                                                                                                                                                                                                                                                   WPI; 2005-112720/12
                                                                                                                                                                                                                                                                                                                                                    Blenis J,
                                                                                                                                                                                                                                                                                                                                                                                                                                         (HARD ) HARVARD COLLEGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-JUL-2003; 2003US-0484761P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-JUL-2004; 2004WO-US021514.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2005007090-A2
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                                                                                                                                                                                                                                                                                                                                               Murphy LO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MAP kinase; cytostatic; cardiovascular-gen.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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    Mismatches

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel method for the identification of a CC therapeutic compound. The method involves providing test cells that CC express a target protein containing a DEF domain and MAP kinase; CC culturing the cells in the presence of growth factor, cytokine, tumor CC promoter or oncogene; contacting the cells with a candidate compound; and CC assessing the binding of the MAP kinase to the DEF domain relative to the binding in the absence of the candidate compound. The invention further CC comprises a method for the identification of a therapeutic compound; a CC method for treatment of cancer, which involves administering a compound CC that inhibits the binding of a MAP kinase to the DEF domain of a target CC protein; and an antibody that specifically binds to phospho-T-325 c-Fos (preferably polyclonal or monoclonal) The novel therapeutic compounds CC have the following activities: cytostatic, cardiovascular-gen., cardiant, CC antiallergic, immunosuppressive, antibacterial, antiallammatory, CC antiallergic, immunosuppressive, antibacterial, antiallammatory, CC antipyretic, uncopathic, antipsoriatic, hepatotropic, antianemic, muscular cgen., thyromimetic, antipsoriatic, hepatotropic, antianemic, muscular cgen., thyromimetic, antipsoriatic, hepatotropic, anticheumatic, cc., cardiovascular disorders, manbolic, hypertensive, annotectic, cancer, cardiovascular disorders, hypothe, annotectic, compound may be used in the treatment cof: cancer, cardiovascular disorders, inflammatory disorders, metabolic disorders, metabolic therapy target peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches
                                                                                          07-AUG-2000; 2000US-0223358P
29-SEP-2000; 2000US-0236827P
01-AUG-2001; 2001US-00920262
                                                                                                                                                                                                                                                                                                                                   complementarity determining region; neuroprotective; antitipsoriatic; immunostimulant; cytostatic; anti-microbial; psoriasis; infection; multiple sclerosis; immune disorder; cardiovascular; malignant disease.
                 Giles-Komar J, Knight DM,
                                                                                                                                                                    07-AUG-2001; 2001WO-US024720
                                                                                                                                                                                                                                                                                                                                                                                       Human; antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                             Anti-Interleukin-12 (IL-12) antibody CDR3 light chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAU76520 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 14; Page 64; 104pp; English
                                                        (CENZ ) CENTOCOR INC.
                                                                                                                                                                                                                                          WO200212500-A2
                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                 neurological disorder
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                           anti-interleukin-12; CDR; light chain; circulatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                33.5%;
                     Peritt D,
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Pred. No. 3.4e+02;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to novel isolated mammalian anti-interleukin-12 (IL-12) antibodies. The antibodies comprise at least one complementarity determining region (CDR) of a heavy or light chain, a heavy chain or light chain variable region, or a heavy chain or light chain constant region. The anti-IL-12 antibodies are useful in methods for diagnosing or treating IL-12 related conditions, e.g. psoriasis or multiple sclerosis. The antibodies are also useful for treating immune, cardiovascular, infectious, malignant or neurological disorders or diseases. The present sequence represents the amino acid sequence of human anti-Interleukin-12 (IL-12) antibody CDR3 light chain
                                                         treating
                                                                          Novel
                                                                                                                                                                                                                                              24-MAY-2001; 2001WO-US016864.
                                                                                                                                                                                                                                                                              29-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                  immunoglobulin; complementarity determining region; CDR; protozoacide; antiinflammatory; immunosuppressive; anti-HIV; antibacterial; virucide; fungicide; neuroprotective; light chain variable region; VK; mAb;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; sialoadhesin factor-3; SAF-3; therapy; cancer; inflammation; autoimmunity; allergy; asthma; infection; central nervous system; C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAE15818 standard; peptide; 9
                                                                                                                                       Abrahamson
                                                                                                                                                                                                                 24-MAY-2000; 2000US-00577930.
                                                                                                                                                                                                                                                                                                           WO200190193-A1
                                                                                                                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                                                                                                                      monoclonal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                  rheumatoid arthritis; multiple sclerosis; stem cell mobilisation; AIDS; haematopoietic development; anaemia; chemoprotective agent; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                             autoimmunity; allergy; asthma; infection; central nervous system; rheumatoid arthritis; multiple sclerosis; stem cell mobilisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAE15818
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                                                                                                                                                                    (SMIK)
                                                                                                                                                                                  (SMIK)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mammalian anti-IL-12 antibodies, useful for diagnosing or treating related conditions, e.g. psoriasis or multiple sclerosis, as well a refer treating immune, infectious, malignant or neurological
                                                                                                                                                                                                                                                                                                                                      sapiens
                                                        monoclonal antibody that binds to human sialoadhesin factor-3 for ing or preventing cancer, inflammation, autoimmunity, allergy,
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                                           rheumatoid
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                                                                                                                                     Kikly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            complementarity determining region
                                           arthritis, multiple sclerosis,
                                                                                                                                                                    BEECHAM
                                                                                                                                                                                  BEECHAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           96pp;
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                                                                                                                                                                     PLC
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.1e+06;
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                                           infections
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Claim 13;

Page

67; 69pp;

English

recovering from the sub-library members

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RESULT 12
AEF76312
ID AEF76
XX AEF76
AC AEF76
AC AEF76
AC AEF76
XX DT 20-AP
XX antib
KW neopl
XX neopl
XX neopl
XX neopl
XX NO NO
XX NO
PN WO200
XX 16-FE
XX DOWN
XX NO
PR 10-JU
XX AEF76
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PR 10-JU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 5
                           The invention relates to an antibody (I) that binds to a prostate cancer cell. Also described are the following: identifying (MI) antibodies specific to cancer cells, generating a phage displayed antibody library using cells collected from subjects immunized with cancer cells, removing members of the library that bind to human red blood cells to generate a
                                                                                                                                                                                                                                                                                             New antibody that binds to a prostate cancer cell, useful cancerous cells in vivo, for killing or ablating prostate vivo, and for delivering a variety of cytotoxic drugs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bowdish KS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antibody; prostate tumor; cytostatic; andrology; genitourinary disease; neoplasm; antibody identification; antibody therapy; light chain.
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                                                                                                                                                                                                                                   Claim 16; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-JUL-2004; 2004US-0586811P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-JUL-2005; 2005WO-US024260
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ALEX-) ALEXION PHARM INC
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                                                                                                                                                                                                                                   48pp;
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Pred. No. 2.1e
1; Mismatches
                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Siva
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. 2.1e+06;
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RESULT 13
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PF 24-M
PR 24-M
PR 24-M
PR 25-M
PR 14-M
PR 25-M
PR 25-M
PR 25-M
PR 14-M

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local
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02-APR-2003;
03-APR-2003;
08-APR-2003;
                                                                                                                                                                                                                                                                                                                                             Chan K, (
Leung FC;
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                                                                                                                                                   New nucleic acid molecule encoding nucleocapsid- or spike-gene protein of a human Severe Acute Respiratory Syndrome (hSARS) virus, useful for diagnosing and treating SARS.
                                                                                                                                                                                                                                                                                   WPI; 2004-737326/72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-APR-2003;
23-APR-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Virucide; Severe Acute Respiratory Syndrome; SARS; vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hSARS virus peptide, SEQ ID 1387.
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2003US-0457730P.
2003US-0459931P.
2003US-0460357P.
2003US-0461265P.
2003US-0462805P.
2003US-0464886P.
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2003US-0470935P.
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; Pred. No. 2.1e
1; Mismatches
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2.1e+06;
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The present invention relates to novel human Severe Acute Respiratory Syndrome (hSARS) viral nucleic acid and protein sequences derived from

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08-APR-2003;
14-APR-2003;
                               New isolated nucleic acid molecule useful for detecting, treating, ameliorating, or preventing the virus causing severe acute respiratelyndrome in humans using a real-time quantitative polymerase chain
                                                                                                                                                                                                                                       23-APR-2003;
05-MAY-2003;
                                                                                                                                                                                                                                                                                                                                 26-MAR-2003;
02-APR-2003;
                                                                                                                                                                                                                                                                                                                                                                      24-MAR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SARS coronavirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            virucide; vaccine; detection; severe acute respiratory syndrome; real-time quantitative polymerase chain reaction; SARS.
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                                                                                                       WPI; 2004-737292/72.
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2003US-0460357P.
2003US-0461265P.
2003US-0462805P.
2003US-0464886P.
2003US-0468139P.
2003US-0468139P.
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54.5%;
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Pred. No.
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26-MAR-2003;
02-APR-2003;
03-APR-2003;
08-APR-2003;
The present invention relates to novel human Severe Acute Respiratory Syndrome (hSARS) viral nucleic acid and protein sequences derived from a hSARS virus having China Center for Type Culture Collection Deposit Accession No. CCTCC-V200303. The hSARS virus, nucleic acid and protein sequences are useful as vaccines for diagnosing or treating SARS. They are also useful in clinical and scientific research applications. The hSARS virus genome (ADT36557) was obtained and the amino acid sequences of all three reading frames were deduced from the complementary strand. ADT37650 is the full-length protein encoded by the first reading frame c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to an isolated nucleic acid molecule consisting essentially of, and/or hybridizes under stringent conditions to a fully defined nucleotide sequence of 16-25 base pairs (bp; SEQ ID NO: 2471-2476), or its complement. The methods and compositions of the present invention are useful for the detection of the virus causing Severe Acute Respiratory Syndrome (SARS) in humans using a real-time quantitative polymerase chain reaction (PCR) assay. They can also be used in treating, ameliorating, managing or preventing SARS. This sequence corresponds to a partial SARS protein sequence from the complementary reading frame 1.
                                                                                                                                                                                                                                                                                        New isolated human severe acute respiratory syndrome useful as vaccine for diagnosing or treating SARS or scientific research applications.
                                                                                                                                                                                                                                                                                                                                                                                        HPI;
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3e+02;
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                                                                                                                                                                                                                                                                                                                    (hSARS) virus, in clinical and
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2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

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              US-10-758-165-10
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US-10-912-994-6
US-10-975-788-3
US-10-975-788-3
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US-10-246-019A-1
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US-10-807-807-1387
US-10-807-807-138-138
US-10-107-532-515
US-10-107-532-515
US-10-908-187-151
US-10-908-187-1527
US-10-908-188-126
US-10-908-5-548A-126
US-10-908-5-548A-126
US-10-808-772-27
US-10-808-772-27
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e 9, Appli
312, Appl
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14, Appl
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2080, Ap
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5915, Ap
6124, Appl
1511, Ap
64, Appl
16, Appl
1527, Ap
126, App
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| •                | •                | •                | •                  | 28.9             |                  | •                | •                | •                | •                |                  | •                |                  | •                 | •                | •                  | •                  | ٠                 |
| 10               | 10               | 9                | 9                  | 9                | 9                | 9                | v                | 18               | 18               | 18               | 17               | 15               | 14                | 12               | 9                  | 9                  | v                 |
| u                | 4,               | 0                | ഗ                  | 4.               | 4                | 4                | 4                | ഗ                | 4.               | 4                | 4                | u                | ഗ                 | w                | თ                  | σ                  | σ                 |
| US-10-691-157-18 | US-10-281-652-18 | US-11-218-813-34 | US-10-482-284A-186 | US-10-160-505-34 | US-10-688-015-34 | US-10-449-379-34 | US-10-160-506-34 | US-10-846-548A-1 | US-10-654-200-29 | US-10-654-200-28 | US-10-306-631-76 | US-10-720-831-20 | US-10-865-478-786 | US-09-813-653-26 | US-11-009-769A-214 | US-11-009-873A-214 | 17-H056-600-TT-SO |
| Sequence         | Sequence         | Sequence         | Sequence           | Sequence         | Sequence         | Sequence         | Sequence         | Sequence         | Sequence         | Sequence         | Sequence         | Sequence         | Sequence          | Sequence         | Sequence           | Sequence           | aprianhae         |
| 18,              | 18,              | 34,              | 186                | 34,              | 34,              | 34,              | 34,              | ۲,               | 29,              | 28,              | 76,              | 20,              | 786               | 26,              | 214                | 214,               | 1.17              |
| Appl             | Appl             | Appl             | , Apr              | Appl             | Appl             | Appl             | Appl             | App 1            | Appl             | Appl             | Appl             | Appl             | , Apr             | Appl             | , App              | , Apr              | 100               |

ALIGNMENTS

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; LENGTH: 18
; TYPE: PRT
; ORGANISM: Felis catus
US-10-758-165-10
                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-758-165-10
                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1
                                                                                                                                                                                                                      Sequence 10, Application US/10758165

Publication No. US20050196816A1

GENERAL INFORMATION:

APPLICANT: Hammerberg, Bruce

TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES

FILE REFERENCE: 5051-661

CURRENT APPLICATION NUMBER: US/10/758,165

CURRENT FILING DATE: 2004-01-16

PRIOR APPLICATION NUMBER: US 60/440,472

PRIOR FILING DATE: 2003-01-16

NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                         SOFTWARE: PatentIn version 3.2 SEQ ID NO 10
                                                   Matches
                                                                  Query Match
Best Local Similarity
                                                   18;
1 VDGQKATNIFPYTAPGKQ 18
                                                   Conservative
                                                                    100.0%;
                                                   0
                                                   Score 97; DB 5;
Pred. No. 6.2e-09;
Mismatches 0;
                                                     0
                                                                                   Length 18
                                                     Indels
                                                   °.
                                                     Gaps
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Oy 1 VDGQKATNIFPYTAPGKQ 18

RESULT 2

US-10-758-165-9

REBULT 2

REPERENCE: 5051-661

FILIC REPERENCE: 5051-661

CURRENT APPLICATION NUMBER: US/10/758,165

CURRENT APPLICATION NUMBER: US/10/758,165

CURRENT APPLICATION NUMBER: US/60/440,472

PRIOR APPLICATION NUMBER: US 60/440,472

PRIOR FILING DATE: 2003-01-16

SOFTWARE: PATENT DI NOS: 16

SOFTWARE: PATENT NO SOFTWARE:
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PRIOR APPLICATION NUMBER: PCT/GB02/005411
PRIOR FILING DATE: 2002-12-02
PRIOR APPLICATION NUMBER: GB0128733.3
PRIOR FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: US 60/334,649
PRIOR FILING DATE: 2001-11-30
NUMBER OF SEQ ID NOS: 41
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 4
US-10-856-118-34
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US-10-758-165-12
                                                                                                                                                                                                 ; OTHER INFORMATION: Flanking sequence US-10-856-118-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
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US-10-758-165-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Skinner, Mike
APPLICANT: Hill, Adrian V.S.
APPLICANT: Hill, Adrian V.S.
APPLICANT: Anderson, Richard
TITLE OF INVENTION: Vaccine
FILE REFERENCE: 3742.1000-000
CURRENT APPLICATION NUMBER: US/10/856,118
CURRENT FILING DATE: 2004-05-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:

APPLICANT: Hammerberg, Bruce
TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR FILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin version 3.2
SEQ ID NO 12
LENGTH: 18
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 34, Application US/10856118 Publication No. US20050025747A1 GENERAL INFORMATION:
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Matches 16; Conserva
                                                                                                                                     Query Match
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Publication No. US20050196816A1
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APPLICANT: Skinner, Mike
                                                                                                                                                                                                                                                     LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                     38.1%;
Local Similarity 55.6%;
tes 5; Conservative
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7 TNIFPYTAP 15
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66.7%;
                                                                                                             Score 37; DB Pred. No. 50;
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Pred. No. 0.061;
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); Mismatches 0;
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                                                                                                                                       DB 5;
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در
                                                                                                                                       Length 15;
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US-10-758-165-11
Sequence 11, App
Publication No.
                                                                                                           RESULT 7
US-09-920-262A-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:epitope peptide OTHER INFORMATION: P5, middle portion of C2 of equine IgE epsilon OTHER INFORMATION: heavy chain
US-10-052-788-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       밁
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; TYPE: PRT
; ORGANISM: Equus caballus
US-10-758-165-11
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Sequence 5, Application US/10052788
Publication No. US20030087314A1
GENERAL INFORMATION:
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                            Sequence 6, Application US/09920262A Publication No. US20030124123A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATI: Hammerberg, Bruce
TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR FILING DATE: 203-01-16
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PATENTIA PATE: 203-01-16
SOFTWARE: PATENTIA PATEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 5
LENGTH: 15
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Best Local
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Gershwin, Laurel J.
APPLICANT: Pettigrew, Howard David
APPLICANT: Kalina, Warren V.
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Epsilon Immunoglobulin Chain Derived
TITLE OF INVENTION: Induction of Anti-IgE Antibodies
FILE REFERENCE: 023070-121000US
APPLICANT: Shealy, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/052,788
CURRENT FILLING DATE: 2001-11-08
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn Ver: 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
                                                                                                                                                                                                                                                                            1 VDGQKATNIFP 11
                                                                                                                                                                                                                                                                                                                                   6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
8; Conserv
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vo. US20050196816A1
                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            36.1%; Score 35; DB 5; L6 44.4%; Pred. No. 1.3e+02; ative 1; Mismatches 9;
                                                                                                                                                                                                                                                                                                                                                                35.1%;
                                                                                                                                                                                                                                                                                                                                                             Score 34; DB 4;
Pred. No. 1.6e+02;
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APPLICANT: Giles-Komar, Jill
APPLICANT: Knight, David
APPLICANT: Knight, David
APPLICANT: Peritt, David
APPLICANT: Scallon, Bernie
APPLICANT: Scallon, Bernie
APPLICANT: Shealy, David
TITLE OF INVENTION: ANTI-L1-12 ANTIBODIES, CC
FILE REFERENCE: CEN0248DIV1
CURRENT APPLICATION NUMBER: US/10/912,994
CURRENT FILING DATE: 2004-08-06
PRIOR APPLICATION NUMBER: US 60/223,358
PRIOR FILING DATE: 2000-08-07
PRIOR FILING DATE: 2000-08-07
PRIOR APPLICATION NUMBER: US 60/236,827
PRIOR FILING DATE: 2000-09-29
PRIOR FILING DATE: 2001-08-01
NUMBER OF SEQ ID NOS: 15
COCTUADE: BEG ID NOS: 15
                                                                                                            US-10-975-883-6
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                                                                                                                                  RESULT 9
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US-10-912-994-6
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Sequence 6, Application US/10975883
Publication No. US20050111127A1
GENERAL INFORMATION:
APPLICANT: Giles-Komar, Jill
APPLICANT: Knight, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver 3.1 SEQ ID NO 6
                                                                                                                                                                                                                                                                      unery Match 33.0%;
Best Local Similarity 83.3%;
Matches 5; Conservative
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Publication No. US20050002937A1
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Matches 5; Conserva
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APPLICANT: Giles Komar, Jill
APPLICANT: Giles Komar, Jill
APPLICANT: Peritt, David
TITLE OF INVENTION: ANTI-IL-12 ANTIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CEN0248
CURRENT FILING DATE: 2002-05-06
CURRENT FILING DATE: 2002-05-06
PRIOR APPLICATION NUMBER: 60/223,358
PRIOR FILING DATE: 2000-08-07
PRIOR PPLICATION NUMBER: 60/236,827
PRIOR PPLICATION NUMBER: 60/236,827
PRIOR PILING DATE: 2000-09-29
NUMBER OF SEQ ID NOS: 15
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TYPE: PRT
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TYPE: PRT
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83.3%;
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Pred. No. 1.9e+06;
1; Mismatches 0
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Pred. No. 1.9e+06
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CURRENT APPLICATION NUMBER: US/10/975,883
CURRENT FILING DATE: 2004-10-28
PRIOR APPLICATION NUMBER: US 60/223,358
PRIOR FILING DATE: 2000-08-07
PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: US 60/236,827
PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: US 09/920,262
PRIOR FILING DATE: 2001-08-01
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PATENTIN Ver 3.1
SEQ ID NO 6
                                                                                                                                                                                                                                                                        APPLICANT: Scallon, Bernie
APPLICANT: Shealy, David
TITLE OF INVENTION: ANTI-IL-12 ANTIBODIES AND NUCLEIC ACIDS
FILE REFERENCE: CENO248DIV03
CURRENT APPLICATION NUMBER: US/10/975,740A
CURRENT FILING DATE: 2004-01-28
PRIOR APPLICATION NUMBER: US 60/223,358
PRIOR FILING DATE: 2000-08-07
PRIOR FILING DATE: 2000-08-07
PRIOR APPLICATION NUMBER: US 60/236,827
PRIOR APPLICATION NUMBER: US 60/236,827
PRIOR FILING DATE: 2000-09-29
PRIOR FILING DATE: 2000-09-29
PRIOR FILING DATE: 2000-09-29
PRIOR FILING DATE: 2001-08-01
NUMBER OF SEQ ID NOS: 9
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APPLICANT: Giles-Komar, Jill
APPLICANT: Knight, David
APPLICANT: Peritt, David
APPLICANT: Scallon, Bernie
APPLICANT: Shealy, David
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US-10-975-740A-6
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US-10-975-740A-6
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SEQ ID NO 6
                                                                             Matches
                                                                                            Query Match
Best Local
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TYPE: PRT
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APPLICANT: Shealy, David
TITLE OF INVENTION: ANTI-IL-12 ANTIBODIES AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens
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                                                                                                                                                                                              TYPE: PRT
                                                                                                                                                                                                                      ENGTH:
4 NIYPYT 9
                                     8 NIFPYT 13
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                                                                                              Similarity
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                                                                             Conservative
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83.3%;
                                                                                              33.0%;
                                                                           Score 32; DB 5;
Pred. No. 1.9e+06;
1; Mismatches 0
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Pred. No. 1.9e+06;
1; Mismatches 0;
                                                                                                                 Length 9;
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RESULT 11 US-10-975-708-6

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Sequence 6, Application US/10975708
Publication No. US20050214293A1
GENERAL INFORMATION:
APPLICANT: Giles-Komar, Jill

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APPLICANT: Kikly, Kristine K.

APPLICANT: Abzahmson, Julie

ITITLE OF INVENTION: Sialoadhesin Factor-3 Antibodies

FILE REFERENCE: GH50019-1C1

CURRENT APPLICATION NUMBER: US/10/246,019A

CURRENT FILING DATE: 2002-09-18

PRIOR APPLICATION NUMBER: USSN 09/577,930

PRIOR APPLICATION NUMBER: USSN 09/046,736

PRIOR FILING DATE: 1998-03-24

PRIOR APPLICATION NUMBER: USSN 60/041,885

PRIOR FILING DATE: 1997-04-02

NUMBER OF SEG ID NOS: 14

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 14
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US-10-246-019A-14
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-975-708-6
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APPLICANT: Scallon, Bernie
APPLICANT: Shealy, David
TITLE OF INVENTION: ANTI-IL-12 ANTIBODIES ANI
FILE REFERENCE: CEN0248DIV02
CURRENT APPLICATION NUMBER: US/10/975,708
CURRENT FILING DATE: 2004-10-28
PRIOR APPLICATION NUMBER: US 60/233,358
PRIOR APPLICATION NUMBER: US 60/236,827
PRIOR FILING DATE: 2000-09-29
PRIOR FILING DATE: 2000-09-29
PRIOR FILING DATE: 2001-08-01
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver 3.1
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Sequence 1387, Application US/10808187
Publication No. US20050009009A1
GENERAL INFORMATION:
APPLICANT: PEIRIS, JOSEPH S. M.
APPLICANT: YUEN, KWOK YUNG
APPLICANT: POON, LIT MAN
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Best Local (
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Best Local Similarity 83.3%;
Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 9
TYPE: PRT
ORGANISM: Human
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83.3%;
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Pred. No. 1.9e+06,
1; Mismatches (
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Pred. No. 1.9e+06
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APPLICANT: NICHOLLS, JOHN M.
APPLICANT: NICHOLLS, JOHN M.
TITLE OF INVENTION: A HIGH-THROUGHPUT DIAGNOSTIC ASSAY FOR THE HUMAN VIRUS
FILE OF INVENTION: CAUSING SEVERE ACUTE RESPIRATORY SYNDROME (SARS)
FILE REFERENCE: V9661.0077
CURRENT APPLICATION NUMBER: US/10/807,807
CURRENT FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/457,031
PRIOR FILING DATE: 2003-03-24
PRIOR FILING DATE: 2003-03-24
PRIOR FILING DATE: 2003-03-24
PRIOR FILING DATE: 2003-03-26
PRIOR FILING DATE: 2003-04-03
PRIOR FILING DATE: 2003-04-04
PRIOR FILING DATE: 2003-04-07
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US-10-807-807-1387
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PRIOR FILING DATE: 2003-04-08
PRIOR PELICATION NUMBER: 60/462,805
PRIOR APPLICATION NUMBER: 60/468,139
PRIOR APPLICATION NUMBER: 60/464,139
PRIOR FILING DATE: 2003-05-05
PRIOR APPLICATION NUMBER: 60/464,886
PRIOR APPLICATION NUMBER: 60/471,200
PRIOR FILING DATE: 2003-05-16
PRIOR PRICING DATE: 2003-05-16
NUMBER: 05/471,200
PRIOR FILING DATE: 2003-05-16
NUMBER: 05/471,200
PRIOR FILING DATE: 2003-05-16
NUMBER: 05/471,200
PRIOR FILING DATE: 2003-05-16
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SEQ ID NO 1387
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Best Local
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APPLICATT: NICHOLLS, JOHN

TITLE OF INVENTION: A DIAGNOSTIC ASSAY FOR THE HUMAN VIRUS CAUSING SEVERE /
TITLE OF INVENTION: RESPIRATORY SYNDROME (SARS)

FILE REFERENCE: V9661.0078

FILE REFERENCE: V9661.0078

CURRENT FILING DATE: 2004-03-24

PRIOR APPLICATION NUMBER: 60/457,031

PRIOR FILING DATE: 2003-03-24

PRIOR APPLICATION NUMBER: 60/457,730

PRIOR APPLICATION NUMBER: 60/457,730

PRIOR APPLICATION NUMBER: 60/459,931

PRIOR APPLICATION NUMBER: 60/459,931

PRIOR APPLICATION NUMBER: 60/460,357

PRIOR PRIOR APPLICATION NUMBER: 60/460,357

PRIOR PRIOR APPLICATION NUMBER: 60/460,357

PRIOR FILING DATE: 2003-04-03

PRIOR FILING DATE: 2003-04-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: LEUNG, FREDERICK C.
APPLICANT: PEIRIS, JOSEPH S. M.
APPLICANT: YUEN, KWOK YUNG
APPLICANT: POON, LIT MAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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APPLICATION NUMBER: 60/464,886
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6; Conservative
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54.5%;
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Pred. No. 2.4e+02
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APPLICANT: Chye, Mee Lee
APPLICANT: Li, Hong Ye
APPLICANT: Ramalingam, Sathiskumar
APPLICANT: Remalingam, Sathiskumar
APPLICANT: Poon, Leo Lit Man
APPLICANT: Peiris, Joseph Sriyal Malik
TITLE OF INVENTION: Genetically Modified Plants Comprising SARS-CoV Viral Nucleotide
TITLE OF INVENTION: Sequences and Methods of Use Thereof For Immunization Against SA
FILE REFERENCE: 2587/73166/RDK
CURRENT FILING DATE: 2004-12-03
PRIOR APPLICATION NUMBER: US/11/004,399
CURRENT FILING DATE: 2004-12-03
PRIOR FILING DATE: 2003-12-03
NUMBER OF SEQ ID NOS: 4043
SOFTWARE: Patentin version 3.1
SEQ ID NO 2080
LENGTH: 11
TYPE: PRT
ORGANISM: SARS-COV Virus
US-11-004-399-2080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; PRIOR FILING DATE: 2003-04-23
; PRIOR APPLICATION NUMBER: 60/465,738
; PRIOR FILING DATE: 2003-04-25
; PRIOR APPLICATION NUMBER: 60/470,935
; PRIOR FILING DATE: 2003-05-14
; NUMBER OF SEQ ID NOS: 2487
; SOCTWARE: Patentin ver: 3.2
; SEQ ID NO 1387
; TYBE: PRT
; ORGANISM: Human severe acute respiratory system virus
US-10-807-807-1387
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US-11-004-399-2080
US-11-004-399-2080
; Sequence 2080, Application US/11004399
; Publication No. US20060053516A1
; GENERAL INFORMATION:
Search completed: May 30, 2006, 15:20:14 Job time : 49.3333 secs
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Best Local Similarity 54.5%;
Matches 6; Conservative
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Local Similarity 54.5%;
hes 6; Conservative
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                                                                                      GOSOSNILPOT 11
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                                                                                                                                                                          Score 32; DB 6; Length 11; Pred. No. 2.4e+02; 1; Mismatches 4; Indels
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1: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US09_NEW_PUB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US10_NEW_PUB.pep:*

6: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US10_NEW_PUB.pep:*

7: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US11_NEW_PUB.pep:*

7: /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
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US-11-219-563-34
US-11-254-182-12
US-11-177-359C-16
US-11-177-359C-16
US-11-177-359C-16
US-11-177-359C-16
US-10-528-066-747
US-09-731-899-19
US-10-528-276-38
US-11-123-959A-83
US-11-124-0-87A-1314
US-11-134-228A-8
US-11-121-28A-79
US-11-121-28A-79
US-11-140-487A-211
US-11-140-487A-211
US-11-140-487A-1346
US-11-129-53-10
US-11-29-563-8
US-11-129-563-8
US-11-129-563-8
US-11-129-563-8
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                                                                                                   Sequence 34, Appl Sequence 16, Appl Sequence 16, Appl Sequence 16, Appl Sequence 1747, Appl Sequence 38, Appl Sequence 38, Appl Sequence 31, Appl Sequence 82, Appl Sequence 82, Appl Sequence 87, Appl Sequence 87, Appl Sequence 87, Appl Sequence 21, Appl Sequence 21, Appl Sequence 318, Appl Sequence 3188, Appl Sequen
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812, App
59, Appl
52, Appl
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| 22 22.7 18 7 US-11-257-998-33 21 21.6 9 7 US-11-140-487A-120 21 21.6 9 7 US-11-140-487A-120 21 21.6 9 7 US-11-140-487A-1446 21 21.6 9 7 US-11-140-487A-1446 21 21.6 10 7 US-11-140-487A-1447 21 21.6 10 7 US-11-140-487A-1596 21 21.6 11 6 US-10-540-431-11 21 21.6 11 7 US-11-177-358C-3 21 21.6 11 7 US-11-177-359C-3 21 21.6 11 7 US-11-177-359C-3 21 21.6 11 7 US-11-177-340C-3 21 21.6 11 7 US-11-177-340C-3 21 21.6 11 7 US-11-177-340C-3 21 21.6 12 7 US-11-177-340C-3 21 21.6 13 7 US-11-177-340C-3 21 21.6 13 7 US-11-175-021 21 21.6 12 10S-09-784-950-21 21 21.6 13 7 US-11-140-487A-8 20 20.6 9 7 US-11-140-487A-212 20 20.6 9 7 US-11-140-487A-212 20 20.6 9 7 US-11-140-487A-212 | 45                 | 44       | 43                | 42          | 41              | 40                | 39               | 38       | 37       | 36            | 35            | 34       | 33       | 32       | 31       | 30    | 29    | 28       | 27        | 26        |
|--|--------------------|----------|-------------------|-------------|-----------------|-------------------|------------------|----------|----------|---------------|---------------|----------|----------|----------|----------|-------|-------|----------|-----------|-----------|
| 10 7 US-11-10-10-10-10-10-10-10-10-10-10-10-10-  | 20                 | 20       | 20                | 20          | 21              | 21                | 21               | 21       | 21       | 21            | 21            | 21       | 21       | 21       | 21       | 21    | 21    | 21       | 21        | 22        |
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|  | 7                  | 7        | 7                 | 7           | 7               | 7                 | _                | 7        | 7        | 7             | 7             | σ        | 7        | 7        | 7        | 7     | 7     | 7        | 7         | ,         |
|  | US-11-140-487A-327 | -11-     | US-11-140-487A-24 | 140-487A-   | US-11-211-175-1 | US-11-122-986-656 | US-09-784-950-21 | -11      | 1        | -11-177-359C- | -11-177-358C- | ٠.       | -11-3    | -11      | -11      | -11-1 | -11-1 | -11-     | -11-055-0 | 6-/C7-TT- |
|  | 327, App           | 212, App | Þ                 | <b>&gt;</b> | 1, Appli        | 656, App          |                  | 800, App | 3, Appli | 3, Appli      | •             | 11, App1 | 1596, Ap | 1447, Ap | 223, App | .0    |       | 120, App | -         | 7         |

ALIGNMENTS

## TITLE OF INVENTION: MODIFIED ANTIBODIES TO PROSTATE-SPECIFIC TITLE OF INVENTION: MEMBRANE ANTIGEN AND USES THEREOF FILE REFERENCE: 13651.001 (BZL-001) CURRENT APPLICATION UNWBER: US/11/219,563 CURRENT FILING DATE: 2005-09-02 PRIOR APPLICATION NUMBER: PCT/US04/06586 PRIOR FILING DATE: 2004-03-03 PRIOR APPLICATION NUMBER: PCT/US04/06586 PRIOR FILING DATE: 2003-03-03 PRIOR APPLICATION NUMBER: US 10/379,838 PRIOR APPLICATION NUMBER: US 10/379,838 PRIOR APPLICATION NUMBER: 10/449,379 PRIOR FILING DATE: 2003-05-30 NUMBER OF SEO ID NOS: 144 RESULT 1 US-11-219-563-34 ; Sequence 34, Application US/11219563 ; Publication No. US20060088539A1 ; GENERAL INFORMATION: ; APPLICANT: Bander, Neil 밁 Ś LENGTH: 9 ; TYPE: PRT ; ORGANISM: Mus musculus US-11-219-563-34 Query Match Best Local Similarity 63.8 Marches 7; Conservative SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 34 1 ĠQSYT--FPYT 9 GOKATNIFPYT 13 28.9%; 0 Score 28; DB 7; Pred. No. 5.2e+04; Mismatches Length 9; 2 Gaps

1:

APPLICANT: ANDYA, JAMES
APPLICANT: GWEE, SHIANG C.
APPLICANT: LIU, JUN
APPLICANT: SHEN, YE
APPLICANT: SHEN, YE
TITLE OF INVENTION: ANTIBODY FORMULATIONS
FILE REFERENCE: P2104R1
CURRENT APPLICATION NUMBER: US/11/254,182
CURRENT FILING DATE: 2005-10-19
PRIOR APPLICATION NUMBER: US 60/620,413

RESULT 2 US-11-254-182-12

Sequence 12, Application US/11254182 Publication No. US20060088523A1 GENERAL INFORMATION:

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8
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                                                                                                                        Sequence 16, Application US/11177359C

Publication No. US20060094058A1

GENERAL INFORMATION:
APPLICANT: CAMPBELL, DOUGLAS A.

TITLE OF INVENTION: PEPTIDE SEQUENCE TAGS AND METHOD OF USING SAME
FILE REFERENCE: 113703-1004

CURRENT FILING DATE: US/11/177,359C

CURRENT FILING DATE: 2005-07-11

NUMBER OF SEQ ID NOS: 16

SOFTWARE: PATENTION: DECEMBER OF SET INCOME.

TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
COMPANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 16, Application US/11177358C

Publication No. US20060094072A1

GENERAL INFORMATION:
APPLICANT: CAMPBELL, DOUGLAS A.

TITLE OF INVENTION: PEPTIDE SEQUENCE TAGS AND METHOD OF IFILE REFERENCE: 113703-1006

CURRENT APPLICATION UNMBER: US/11/177,358C

CURRENT FILING DATE: 2005-07-11

NUMBER OF SEQ ID NOS: 16

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 16

LENGTH: 12

TYPE: PRT

ORGANISM: Artificial Sequence
FEATURE:
FEATURE: PRT

ORGANISM: Artificial Sequence
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US-11-177-358C-16
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; OTHER INFORMATION:
US-11-254-182-12
                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic peptide US-11-177-359C-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic peptide
US-11-177-358C-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2004-10-20
NUMBER OF SEQ ID NOS: 74
SEQ ID NO 12
                                               Query Match
Best Local (
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Best Local Similarity 50.0%;
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ORGANISM: Artificial sequence
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                              Similarity 50.4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                               25.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26.8%;
                              Score 25; DB : Pred. No. 94; 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 25; DB
Pred. No. 94;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Mismatches
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Pred.
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5.2e+04;
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                                                               Length 12;
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APPLICANT: Epimmune Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7,
TITLE OF INVENTION: Peptides and Compositions
FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 767
SOFTWARE: PATENTIN Version 3.2
SOFUMARE: Patentin version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; LENGTH: 12
; TYPE: PRT
; ORGANISM: Artificial Sequence
; PEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-11-177-340C-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 4; Conserva
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Sequence 747, Application US/10;

Publication No. US20060094649A1

GENERAL INFORMATION:
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US-09-731-899-19
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                                                    Sequence 19, Application US/09731899 Publication No. US20060088548A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/11/177,340C
CURRENT FILING DATE: 2005-07-11
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 16
LENGTH: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 16, Application US/11177340C Publication No. US20060099663A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                      Matches
APPLICANT: Chain, Benjamin
TITLE OF INVENTION: CHIMERIC PEPTIDES
TITLE OF INVENTION: FOR IMMUNIZATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: CAMPBELL, DOUGLAS A.
TITLE OF INVENTION: PEPTIDE SEQUENCE TAGS AND METHOD OF USING SAME
FILE REFERENCE: 113703-1005
                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: A3 non-natural consensus peptide
                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
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                                                                                                                                                                                                                                                                       Similarity
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Pred. No.
                                                                                                                                                                                                                                                                       Score 24;
Pred. No.
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AS IMMUNOGENS, USING CHIMERIC
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ANTIBODIES THERETO, AND METHODS PEPTIDES OR ANTIBODIES
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FILE REFERENCE: 20555/1203433-US1
CURRENT APPLICATION NUMBER: US/09/731,899
CURRENT FILING DATE: 2000-12-08
PRIOR APPLICATION NUMBER: 60/169,687
PRIOR PTILING DATE: 1999-12-08
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.3
SEQ ID NO 19
LENGTH: 16
TYPE: PAT
ORGANISM: Tetanus toxin bacteria
US-09-731-899-19
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Sequence 38, Application US/11252276

| Sequence 38, Application US/2006094868A1
| Publication No. US2006094868A1
| GENERAL INFORMATION:
| ApplicanT: Bright, Gary
| ApplicanT: Burroughs-Tencza, Sarah
| ApplicanT: Burroughs-Tencza, Sarah
| ApplicanT: Do INVENTION: A System for Cell Based Screening
| TITLE OF INVENTION NUMBER: US/11/252,276
| CURRENT APPLICATION NUMBER: US/11/252,276
| CURRENT FILING DATE: 2005-10-17
| CURRENT FILING DATE: 2005-10-17
| PRIOR APPLICATION NUMBER: 09/713,572
| PRIOR APPLICATION NUMBER: 09/713,572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 4***
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PEATURE:
OTHER INFORMATION: Description of Unknown Sequence: Haemaglutinin OTHER INFORMATION: epitope from unknown organism US-10-522-356-3
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Publication No. US20060105323A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: WHITELAW, CHRISTOPHER BRUCE ALEXANDER
APPLICANT: WHECK, CHARLES ROLAND
APPLICANT: WOLF, CHARLES ROLAND
APPLICANT: WOLF, CHARLES ROLAND
APPLICANT: WOLF, CHARLES ROLAND
APPLICANT: WOLF, CHARLES ROLAND
TITLE OF INVENTION: WULTI-REPORTER GENE WODEL FOR TOXICOLOGICAL SCREENING
TILLE REFERENCE: 102286.155 US,
TILLE REPERENCE: 102286.155 US,
TOXICOLOGICAL SCREENING
TILLING DATE: 2005-01-26
PRIOR PILLING DATE: 2003-07-25
PRIOR FILLING DATE: 2003-07-25
PRIOR FILLING DATE: 2003-07-26
PRIOR FILLING DATE: 2003-07-26
PRIOR FILLING DATE: 2003-07-26
PRIOR FILLING DATE: 2003-07-26
PRIOR PILLING DATE: 2003-07-26
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US-11-252-276-38
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Best Local S
Matches 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENGTH: 9
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3; Conserv
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; Pred. No. 1.9e
1; Mismatches
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pred. No. 5.2e+04;
1; Mismatches 2
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1.9e+02;
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PRIOR APPLICATION NUMBER: 09/398,965
PRIOR FILING DATE: 1999-09-17
PRIOR PILING DATE: 1999-09-17
PRIOR FILING DATE: 1999-09-17
PRIOR FILING DATE: 1999-09-17
PRIOR FILING DATE: 1998-02-27
PRIOR APPLICATION NUMBER: 09/031,271
PRIOR APPLICATION NUMBER: 08/810,983
PRIOR PILING DATE: 1997-02-27
PRIOR FILING DATE: 1997-02-27
PRIOR FILING DATE: 1997-02-27
PRIOR FILING DATE: 1999-05-26
PRIOR FILING DATE: 1999-05-26
PRIOR APPLICATION NUMBER: 60/106,308
PRIOR FILING DATE: 1998-10-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/11/023,959A
CURRENT FILING DATE: 2004-12-28
CURRENT FILING DATE: 2004-12-28
PRIOR APPLICATION NUMBER: PCT/GB03/002804
PRIOR FILING DATE: 2003-06-30
PRIOR FILING DATE: 2003-06-30
PRIOR FILING DATE: 1003-06-30
PRIOR FILING DATE: 2003-06-30
SEQ ID NO 83
SEQ ID NO 83
SEQ ID NO 83
LENGTH: 9
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US-11-023-959A-83
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Best Local 9
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US-11-140-487A-1314
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TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
                                                     Query Match
Best Local Similarity 50...
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Artificial sequence
CURRENT APPLICATION NUMBER: US/11/140,487A CURRENT FILING DATE: 2005-05-31 PRIOR APPLICATION NUMBER: EP 04012951.2
                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE: 1999-10
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Pred. No. 5.2e+04;
Pred. No. 5.2e+04; Indels
                                                                                                                                                                                                                                                                                                                                         1; Mismatches
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Pred. No. 5.2e+04;
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APPLICANT: SITVABLAVA, FILTER
APPLICANT: CHANDAWARKER, Raji
FITTLE OF INVENTION: METHODS TO IDENTIFY COMPOUNDS USEFUL FOR TITLE OF INVENTION: METHODS TO IDENTIFY COMPOUNDS USEFUL FOR TITLE OF INVENTION: OF PROLIFERATIVE AND DIFFERENTIATIVE DISC CURRENT APPLICATION NUMBER: US/11/106,014
PRIOR FILING DATE: 2005-04-13
PRIOR PILING DATE: 2005-04-31
PRIOR PILING DATE: 2003-07-30
PRIOR APPLICATION NUMBER: 10/632,150
PRIOR APPLICATION NUMBER: 09/385,219
PRIOR APPLICATION NUMBER: 09/385,219
PRIOR FILING DATE: 1999-08-27
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: 60/098,355
PRIOR FILING DATE: 1998-08-28
PRIOR APPLICATION NUMBER: 60/098,355
PRIOR APPLICATION NUMBE
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                                                                                                                                    Matches
                                                                                                                                                                           Query Match
Best Local
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PRIOR APPLICATION NUMBER: EP 04447239.7
PRIOR FILING DATE: 2004-10-28
PRIOR PRIOR FILING DATE: 2005-03-25
PRIOR APPLICATION NUMBER: EP 05102441.2
PRIOR FILING DATE: 2005-03-25
PRIOR APPLICATION NUMBER: US 60/576,310
PRIOR APPLICATION NUMBER: US 60/622,782
PRIOR FILING DATE: 2004-106-03
PRIOR FILING DATE: 2004-10-29
PRIOR PRIOR APPLICATION NUMBER: US 60/665,395
PRIOR APPLICATION NUMBER: US 60/665,395
NUMBER OF SEQ ID NOS: 2278
SEQ ID NO 1314
I.FNCTH. 10
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APPLICANT: Pagano, Michele
APPLICANT: Chiaur, Dah Sha
APPLICANT: Latres, Esther
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: hepatitis C virus US-11-140-487A-1314
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Best Local Similarity 30.0%;
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Srivastava, Promod
APPLICANT: Chandawarker, Rajiv
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                                                    2 DGQKATNIFPYTA 14
                                                                                                                   Similarity 46.2
6; Conservative
DGEKDT--YSYLA 12
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No. US20060088846A1
                                                                                                                                               23.7%;
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                                                                                          Score 23; DB 7; Length 12;
Pred. No. 2.1e+02;
2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 23; DB 7;
Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                         Length 12;
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3 DISORDERS
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RESULT 13

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HARMACEUTICALS INC.

APPLICANT: AUGNTIS PHARMACEUTICALS INC.

APPLICANT: GUO, Yong

APPLICANT: WAO, Zhengbin

TITLE OF INVENTION: MEMBRANE PENETRATING PEPTIDES AND USES THEREOF

CURRENT APPLICATION NUMBER: US/11/251,734

PRIOR APPLICATION NUMBER: US 09/933,780

PRIOR PILING DATE: 2001-08-21

PRIOR FILING DATE: 2001-08-21

PRIOR APPLICATION NUMBER: US 60/227,647

PRIOR APPLICATION NUMBER: US 60/227,647

PRIOR PRILING DATE: 2001-08-25

PRIOR APPLICATION NUMBER: US 60/227,647

PRIOR PRILING DATE: 2001-08-25

PRIOR APPLICATION NUMBER: GB 0103110.3

NUMBER OF SEQ ID NOS: 54

SEQ ID NO 30

IUMBER OF SEQ ID NOS: 54

SEQ ID NO 30

ORGANISM: Artificial Sequence

PEACURE:

OTHER INFORMATION: Synthetic peptide
RESULT 15
US-11-122-986-799
; Sequence 799, Application US/11122986
; Publication No. US20060104989A1
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                                                                                                                                                 Query Match
Best Local Similarity 66.
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: Artificial Sequence; FEATURE; OTHER INFORMATION: Description of Artificial Sequence: Hemagglutinin; OTHER INFORMATION: epitope
US-11-134-228A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 30, Application No. US20, GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 23.7
Best Local Similarity 50.0
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-11-134-228A-8
                                                                                               2 KKTNLF 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US20060100134A1
                                                                                                                                                           23.7%; Score 23; DB 7; 66.7%; Pred. No. 3.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23.7%;
                                                                                                                                               1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 23; DB 7; Leary-Pred. No. 2.3e+02;
                                                                                                                                                                      Length 17;
                                                                                                                                         0,
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                                                                                                                                    Gaps
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                                                                                                                                      0,
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GENERAL INFORMATION:
APPLICANT: EDWARNS; ALED
APPLICANT: DWARNS; ALED
APPLICANT: DWARNS; ALED
APPLICANT: VEDANI, MASOUD
ITIZE OF INVENTION: ESSENTIAL NOVEL BACTERIAL POLYPEPTIDES
ITIZE OF INVENTION: ESSENTIAL STATE
ITIZE OF INVENTION: ESSENTIAL STATE
CURRENT FILLING DATE: 2002-11-05
PRIOR PILLING DATE: 2002-11-06
PRIOR P
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GenCore version 5.1.8
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Database
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Maximum Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         seq
 protein search, using sw model
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Match
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Gapop 10.0 ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         May 30, 2006, 15:08:01; Search time 19.3333 Seconds (without alignments) 81.494 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-758-165A-9
97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                650591 seqs, 87530628 residues
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VDGQKATNIFPYTAPGTK 18
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           US-09-920-262A-6
US-09-674-973A-64
US-09-865-548A-126
US-08-333-565-22
US-08-661-479-26
US-08-758-417A-214
US-09-307-265A-14
US-09-307-265A-14
US-09-526-195-9
US-09-641-803-18
US-09-641-803-18
US-09-641-801-18
US-09-641-801-18
US-09-641-801-28
US-09-641-801-28
US-09-641-801-18
US-09-641-801-28
US-09-641-801-18
US-09-641-801-28
US-09-641-801-28
US-09-641-801-28
US-09-641-801-28
US-09-641-801-28
US-09-641-801-32
US-09-98-322-83
US-09-98-322-83
US-09-98-322-83
US-09-98-322-845-5
US-07-963-329A-69
                                                                                                                                                                                                                                                                                                                                                                                                                          SUMMARIES
             Sequence 6, Appli
Sequence 126, Appl
Sequence 22, Appl
Sequence 22, Appl
Sequence 214, Appl
Sequence 214, Appl
Sequence 214, Appl
Sequence 20, Appli
Sequence 10, Appli
Sequence 11, Appli
Sequence 11, Appli
Sequence 11, Appli
Sequence 18, Appli
Sequence 26, Appli
Sequence 31, Appli
Sequence 41, Appli
Sequence 5, Appli
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| 45              | 44                   | 43                   | 42               | 41               | 40               | 39               | 38               | 37              | 36                 | 35              | 34              | 33                | 32              | 31              | 30               | 29                 | 28                |             |
|-----------------|----------------------|----------------------|------------------|------------------|------------------|------------------|------------------|-----------------|--------------------|-----------------|-----------------|-------------------|-----------------|-----------------|------------------|--------------------|-------------------|-------------|
| 26              | 26.5                 | 26.5                 | 27               | 27               | 27               | 27               | 27               | 27              | 27                 | 27              | 27              | 27                | 27              | 27              | 27               | 27                 | 27                | ,           |
| 26.8            | 27.3                 | 27.3                 | 27.8             | 27.8             | 27.8             | 27.8             | 27.8             | 27.8            | 27.8               | 27.8            | 27.8            | 27.8              | 27.8            | 27.8            | 27.8             | 27.8               | 27.8              |             |
| თ               | 15                   | 15                   | 18               | 18               | 17               | 17               | 17               | 17              | 13                 | 13              | 13              | 13                | 13              | 13              | 13               | 11                 | 11                | ;           |
| N               | w                    | w                    | <b>_</b>         | ۳,               | N                | N                | ۳                | ۳               | IJ                 | N               | N               | N                 | ۲               | ۳               | ۲                | v                  | N                 | 1           |
| US-09-788-006-9 | US-09-641-528B-47461 | US-09-641-528B-47440 | US-09-017-205-49 | US-09-017-205-48 | US-08-466-381C-9 | US-08-705-477E-9 | US-08-394-152A-9 | US-08-325-553-9 | PCT-US92-09443A-58 | US-09-064-159-8 | US-09-318-001-8 | US-07-963-329A-58 | US-08-823-245-8 | US-08-462-018-8 | US-07-958-903A-8 | PCT-US92-09443A-69 | US-09-809-917A-21 |             |
| Sequence        | Sequence             | Sequence             | Sequence         | Sequence         | Sequence         | Sequence         | Sequence         | Sequence        | Sequence           | Sequence        | Sequence        | Sequence          | Sequence        | Sequence        | Sequence         | Sequence           | Sequence          | 1 1 1 1 1 1 |
| 9, Appli        | 47461, A             | 47440, A             | 49, Appl         | 48, App1         | 9, Appli         | 9, Appli         | 9, Appli         | 9, Appli        | 58, Appl           | 8, Appli        | 8, Appli        | 58, Appl          | 8, Appli        | 8, Appli        | 8, Appli         | 69, Appl           | 21, App1          |             |

## ALIGNMENTS

```
GENERAL INFORMATION:
APPLICANT: Shealy, David
APPLICANT: Shealy, David
APPLICANT: Scallon, Bernie
APPLICANT: Scallon, Bernie
APPLICANT: Scallon, Bernie
APPLICANT: Gelies-Komar, Jill
APPLICANT: Peritt, David
ITILE OF INVENTION: ANTI-IL-12 ANTIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CENO248
CURRENT APPLICATION NUMBER: US/09/920,262A
CURRENT APPLICATION NUMBER: US/09/920,262A
CURRENT APPLICATION NUMBER: 60/233,358
PRIOR APPLICATION NUMBER: 60/233,358
PRIOR APPLICATION NUMBER: 60/236,827
PRIOR APPLICATION NUMBER: 60/236,827
PRIOR APPLICATION ONDER: 50/236,827
PRIOR APPLICATION ONDER: 50/236,827
PRIOR FILING DATE: 2000-09-29
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver 3.1
SEQ ID NO 6
LENGTH: 9
Sequence 64, Application US/09674973A
Patent No. 6759046
GENERAL INFORMATION:
APPLICANT: NO. 67590468k Hydro ASA
TITLE OF INVENTION: Peptides
FILE REFERENCE: 26625-296
CURRENT APPLICATION UMMBER: US/09/674,973A
CURRENT FILING DATE: 2001-06-04
NUMBER OF SEQ ID NOS: 459
SOFTWARE: PatentIn version 3.0
SEQ ID NO 64
LENGTH: 15
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                                                                                                                                                                                                                                                                                                              RESULT 2
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Best Local S
Matches 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                           33.0%;
Local Similarity 83.3%;
tes 5; Conservativo
                                                                                                                                                                                                                                                                                                                                                                               4 NIYPYT 9
                                                                                                                                                                                                                                                                                                                                                                                                                           8 NIFPYT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 32; DB 2;
Pred. No. 5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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RESULT 4
US-08-333-565-22
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US-09-865-548A-126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 22, Application US/08333565
Patent No. 5622852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 126, Application US/09865548A Patent No. 6867283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Admon, Arie
TITLE OF INVENTION: METHOD OF IDENTIFYING PEPTIDES CAPABLE OF BINDING TO MHC MOLECULE
TITLE OF INVENTION: PEPTIDES IDENTIFIED THEREBY AND THEIR USES
FILE REFERENCE: 0/1/22080
FILE REFERENCE: 0/1/22080
CURRENT APPLICATION NUMBER: US/09/865,548A
CURRENT FILING DATE: 2001-05-16
PRIOR APPLICATION NUMBER: US 60/290,958
PRIOR FILING DATE: 2001-05-16
NUMBER OF SEQ ID NOS: 204
SOFTWARE: PATENTIN OF SEQ ID NOS: 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:

APPLICANT: KORSMEYER, Stanley

TITLE OF INVENTION: Bcl-x/Bcl-

TITLE OF INVENTION: REGULATOR

NUMBER OF SEQUENCES: 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Barnea, Eilon
APPLICANT: Beer, Ilan
APPLICANT: Ziv, Tamar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: synthetic peptide
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/333,565
FILING DATE: 31-OCT-1994
CLASSIFICATION: 435
                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       33.0%;
Local Similarity 54.5%;
les 6; Conservarino
                                                                                                                                                                                                                COUNTRY: U
                                                                                                                                                                                                                                                            CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 100 tes 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8 NIFPY 12
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                                                                                                                                                                                                                                                                                                   379 Lytton Avenue
                                                                                                                                                                                                                                        S
                                                                                                                                                                                                                                                                                                                                                                                                                  KORSMEYER, Stanley J.
VENTION: Bcl-x/Bcl-2 ASSOCIATED CELL DEATH
                                                                                                                                                               Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30.9%; Score 30; DB 2; Length 9; 100.0%; Pred. No. 5e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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Pred. No. 1.2e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-661-479-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 5
US-08-661-479-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 1572
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
TELEPAX: (415) 326-2422
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acids
TYPE: amino acids
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 5834209
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: KORSMEYER, Stanley J.
TITLE OF INVENTION: Bcl-x/Bcl-2 ASSOCIATED CELL DEATH
TITLE OF INVENTION: REGULATOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Loc
Matches
Query Match 30.9%;
Best Local Similarity 83.3%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/661,479
FILING DATE: 11-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/333,565
FILING DATE: 31-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M
DECIFERATION NUMBER: 30-323
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                  TELEFAX: (415) 326-2422
INFORMATION FOR SEQ ID NO:
                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                              REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 15726A-000760
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: STREET: STREET: OF THE CITY: Palo Alto CITY: Palo Alto CITATE: California
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o. 5834209
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ilarity 83.3%;
Conservative
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Pred. No. 2.8e+02;
0; Mismatches 1
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  Score 30; DB 1;
Pred. No. 2.8e+02;
0; Mismatches
                                      Length 17;
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      Indels
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RESULT 6
US-09-042-353-366
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-UN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
                                                                    APPLICATION NUMBER: US 08/209,741
FILING DATE: 09-MAR-1994
PRIOR APPLICATION DATA:
APPLICATION UNUMBER: US 08/352,322
FILING DATE: 07-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/544,404
FILING DATE: 10-CCT-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/728,463
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APPLICATION NUMBER: US 08/155,301
FILING DATE: 18-NOV-1993
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APPLICATION NUMBER:
FILING DATE: 26-APR-
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic
TITLE OF INVENTION: Producing
                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                    PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/096,762 FILING DATE: 22-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 0 FILING DATE: 17-DEC-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 1 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                          FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: San Francisco
                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
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Two Embarcadero
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26-APR-1993
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                                                        10-OCT-1996
MBER: WO PCT/US96/16433
10-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transgenic No. 6255458-Human Animals for Producing Heterologous Antibodies
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Center, Eighth Floor
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RESULT 7
US-08-758-417A-214
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Patent No. 6300129
GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 01.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE: 02-DEC-PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
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FILING DATE: 01-DEC-1997
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                                                                                                                                                                    PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/728,463

FILING DATE: 10-0CT-1996

APPLICATION NUMBER: US 08/544,404

FILING DATE: 10-0CT-1995

APPLICATION UMBER: US 08/352,322

FILING DATE: 07-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kay, Robert M.
TITLE OF INVENTION: Transgenic No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Lonberg, Nils
Kay, Robert M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                     APPLICATION NUMBER: US 08/209,741
FILING DATE: 09-MAR-1994
APPLICATION NUMBER: US 08/165,699
FILING DATE: 10-DEC-1993
APPLICATION NUMBER: US 08/161,739
                                                                                                                                                                                                                                                                                                             CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/758,417A FILING DATE: 02-Dec-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: California
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                : 03-DEC-1993
NUMBER: US 08/155,301
: 18-NOV-1993
NUMBER: US 08/096,762
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Pred. No. 5e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              No. 6300129-Human Animals for Heterologous Antibodies
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GENERAL INFORMATION:
APPLICANT: Gu, Trent
APPLICANT: Gu, Trent
APPLICANT: Gu, Trent
APPLICANT: Gu, Trent
APPLICANT: Han, Min
CURRENT APPLICATION NUMBER: US/09/307,265A
CURRENT APPLICATION NUMBER: US/09/307,265A
CURRENT FILLING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 14
LENGTH: 11
TYPE: PRT
ORGANISM: Caenorhabditis elegans
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Best Local Similarity
"hes 5; Conserva
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MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 214:
US-08-758-417A-214
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Best Local Similarity 83.3
Thes 5; Conservative
                    Sequence 20, Application US/08278774

Patent No. 6653450

GENERAL INFORMATION:
APPLICANT: Berg, Richard A
APPLICANT: Toman, David P
APPLICANT: Wallace, Donald
TITLE OF INVENTION: MUTATED RECOMBINANT COLLAGENS
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: COLLAGEN CORDENERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 14, Application US/09307265A Patent No. 6225456
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FILING DATE: 26-APR-1993
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
ATTORNEY/AGENT INFORMATION:
NAME: Serafini, Andrew T.
                                                                                                                                                                                                                                                                               11 PYTAPGTK 18
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STRANDEDNESS: <Unknown>
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E: COLLAGEN CORPORATION 2500 Faber Place
                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29.9%;
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Pred. No. 5e+05;
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Sequence 9, Application US/09526195
Patent No. 6699478
GENERAL INFORMATION:
APPLICANT: Hancock, Gerald E.
APPLICANT: Hancock, Gerald E.
APPLICANT: Hancock, Gerald W.
TITLE OF INVENTION: ENHANCED IMMUNE RESPONS
TITLE OF INVENTION: G. PROTEIN OF RESPIRAT
FILE REFERENCE: 1646-103-004
CURRENT APPLICATION NUMBER: US/09/526,195
CURRENT FILING DATE: 2000-03-15
PRIOR APPLICATION NUMBER: US 60/084,863
PRIOR FILING DATE: 1998-05-08
PRIOR APPLICATION NUMBER: PCT/US98/19656
PRIOR APPLICATION NUMBER: US 60/059,684
PRIOR FILING DATE: 1998-09-17
PRIOR FILING DATE: 1997-09-19
NUMBER OF SEQ ID NOS: 37
NUMBER OF SEQ ID NOS: 37
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                                                                                                                                      US-09-526-195-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; MOLECULE TYPE: peptide US-08-278-774-20
                                                                                                                                                                                                                                SOFTWARE: F. SEQ ID NO 9
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Loc
Matches
                                                       Query Match 29.9%; Score 29; DB 2; Le Best Local Similarity 50.0%; Pred. No. 3.6e+02; Matches 6; Conservative 0; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                     TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                      OTHER INFORMATION: Synthetic Peptide
                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Rafayko, Kathi L
REGISTRATION NUMBER: 36,644
REFERENCE/DOCKET NUMBER: 94
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 354-4642
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/278,774
FILING DATE: 22-JUL-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 
les 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (415) 354-46
TELEFAX: (415) 354-4752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 94303
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7 TNIFPYTAPGTK 18
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                                                                                                                                                                                                                                                                          FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENHANCED IMMUNE RESPONSE TO ATTACHMENT (G) PROTEIN OF RESPIRATORY SYNCYTIAL VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29.9%;
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Pred. No. 3.6e+02;
2; Mismatches 5;
                                                                                             Length 15;
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                                                             Indels
                                                             0
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                                                         Gaps
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TTILASTTPGVK 12

US-08-957-130-19 RESULT 11

Sequence 19, Application Patent No. 6290959

US/08957130

GENERAL INFORMATION:

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                                      ; TYPE: PRT
; ORGANISM: Influenza virus
US-09-856-920-1
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US-09-856-920-1
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; MOLECULE TYPE:
US-08-957-130-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: WU, AUGTON
APPLICANT: SUN, Tung-Tien
APPLICANT: SUN, Tung-Tien
TITLE OF INVENTION: METHOD FOR SCREENING COMPOUN
TITLE OF INVENTION: INHIBITING BACTERIAL ATTACH
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
STREET: 419 Seventh Street N.W., Ste. 300
                                                                                                              GENERAL INFORMATION:
APPLICANT: Yedda Research and Development Co.
APPLICANT: Yedda Research and Development Co.
TITLE OF INVENTION: Peptide-based vaccine for influenza
FILE REFERENCE: 9822 PCT
CURRENT APPLICATION NUMBER: US/09/856,920
CURRENT FILING DATE: 2002-02-19
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
                                                                                                                                                                                                                                                                                             Sequence 1, Application US/09856920 Patent No. 6740325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 7; Conserv
  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (202) 737-3528 INFORMATION FOR SEQ ID NO: 1
                                                                                                LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: BROWDY, ROGET L.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: SUI
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/957,130
FILING DATE:
CLASSIFICATION: 424
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                    QVTNL----APGTK 13
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    29.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 29; DB 2;
Pred. No. 3.8e+02;
    Score 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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    B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
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Length 18;
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; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-641-803-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ś
                                                                                                                 ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-641-802-18
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US-09-641-803-18
                S
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US-09-641-802-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: P. SEQ ID NO 18
LENGTH: 10
                                                   Query Match
Best Local S
Matches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: STANTON, G. John
APPLICANT: STANTON, G. John
APPLICANT: HUGHES, Thomas K.
APPLICANT: HUGHES, Thomas K.
APPLICANT: BOLDOGH, Istvan
TITLE OF INVENTION: USE OF COLOSTRININ, CONSTITUENT PEPTIDES THEREOF, AND
TITLE OF INVENTION: ANALOGS THEREOF AS OXIDATIVE STRESS REGULATORS
FILE REFERENCE: 265,00220101
CURRENT APPLICATION NUMBER: US/09/641,803
CURRENT FILING DATE: 2000-08-17
PRIOR APPLICATION NUMBER: 60/149,310
PRIOR FILING DATE: 1999-08-17
NUMBER OF SEQ ID NOS: 34
CORTWADER: DATE: 1999-08-17
NUMBER OF SEQ ID NOS: 34
                                                                                                                                                                                                                                             SOFTWARE:
SEQ ID NO 18
                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT:
BOL
                                                                                                                                                                                                                                                                                                                                                                                                                                                   vequence 18, Application US/09641802
Patent No. 6852685
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 44.4%; Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:

USE OF COLOSTRIN

TITLE OF INVENTION: TO PROMOTE NEURAL CH

FILE REFERENCE: 265.00240101

CURRENT APPLICATION NUMBER: US/09/641,802

CURRENT FILING DATE: 2000-08-17

PRIOR APPLICATION NUMBER: 60/149,633

PRIOR FILING DATE: 1999-08-17
                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                            TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                               LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Match 28.9%;
Local Similarity 42.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7 TNIFPYTAP
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9 IFPYTAP 15
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                                                                     Similarity
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                                                                                                                                                                                                                                                                PatentIn Ver. 2.1
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                                                     Conservative
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ON: USE OF
                                                                     28.9%;
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                                                                     Score 28; DB 2;
Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 28; DB 2;
Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                        COLOSTRININ, CONSTITUENT PEPTIDES THEREOF, AND ANALOX NEURAL CELL DIFFERENTIATION
                                                      Mismatches
                                                                                     Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 10
                                                      Indels
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                                                      Gaps
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: STANYON, G. John
APPLICANT: HUGHES, Thomas K.
APPLICANT: BOLDOGH, Istvan
APPLICANT: BOLDOGH, Istvan
APPLICANT: GEORGIADES, Jerzy
TITLE OF INVENTION: THEREOF COLOSTRININ, CONSTITUENT PEPTIDES THEREOF, AND ANALOG
TITLE OF INVENTION: THEREOF CYTOKINES
FILE REFERENCE: 265.00230101
CURRENT FILING DATE: 2000-08-17
PRIOR APPLICATION NUMBER: 60/149,311
PRIOR FILING DATE: 1999-08-17
INUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin ver. 2.1
SEQ ID NO 18
LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
FEARTURE: INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: peptide
US-09-641-801-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8
Search completed: May 30, 2006, 15:09:25 Job time: 20.3333 secs
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US-09-641-801-18

; Sequence 18, Application US/09641801

; Patent No. 6903068
                                                                                                                                                                           Query Match 28.9%; Score 28; DB 2; Length 10; Best Local Similarity 42.9%; Pred. No. 3.3e+02; Matches 3; Conservative 3; Mismatches 1; Indels
                                                                                          9 IFPYTAP 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 VYPFTGP 7
                                                                                                                                                                               0
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OM protein - protein search, using sw model
                                                                                         GenCore version 5.1.8 Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Run on: May 30, 2006, 15:04:28; Search time 12 Seconds (without alignments) 144.325 Million cell updates/sec

Title:
Perfect score:
Sequence: US-10-758-165A-9 97 1 VDGQKATNIFPYTAPGTK 18

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters:

3215

Minimum DB seq length: 0 Maximum DB seq length: 18

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR\_80:\*
1: pir1:\*
2: pir2:\*
3: pir3:\*
4: pir4:\*

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

| 29                 | 28                 | 27     | 26                | 25                 | 24                 | 23                 | 22     | 21                 | 20              | 19     | 18                 | 17              | 16            | 15            | 14                 | 13     | 12     | 11      | 10     | 9      | 80         | 7      | ტ                  | ហ         | 4                  | w      | 2      | μ      | Result<br>No. |
|--------------------|--------------------|--------|-------------------|--------------------|--------------------|--------------------|--------|--------------------|-----------------|--------|--------------------|-----------------|---------------|---------------|--------------------|--------|--------|---------|--------|--------|------------|--------|--------------------|-----------|--------------------|--------|--------|--------|---------------|
| 22                 | 22                 | 22     | 22                | 22                 | 22                 | 23                 | 23     | 23                 | 23              | 23     | 23                 | 23              | 23            | 23            | 23                 | 24     | 24     | 24      | 24     | 25     | 25         | 25     | 25                 | 26        | 27                 | 27     | 27     | 28     | Score         |
|                    |                    |        | 2                 | 22.7               | .2                 | 23.7               | 23.7   | ω.                 | 23.7            | ω.     | ·                  | 23.7            | 23.7          | 23.7          | 23.7               | 24.7   | 4.     | 24.7    | 4.     | 25.8   | 25.8       | 5      | 5                  |           | 7.                 | 27.8   | 7.     |        | Query         |
| 17                 | 15                 | 15     | 14                | 13                 | 12                 | 18                 | 18     | 18                 | 16              | 16     | 16                 | 15              | 13            | 13            | 12                 | 18     | 17     | 15      | 14     | 16     | 16         | 15     | 15                 | 18        | 18                 | 17     | 15     | 15     | Length        |
| N                  | N                  | N      | N                 | N                  | N                  | N                  | N      | N                  | N               | N      | N                  | N               | N             | N             | N                  | N      | N      | N       | N      | N      | N          | N      | N                  | N         | N                  | N      | N      | N      | BB            |
| C84063             | PA0087             | PA0056 | PH1306            | S01904             | PQ0786             | C56046             | S20322 | PQ0072             | S38292          | PS0383 | S03532             | B45115          | C53275        | S23640        | C20907             | A25941 | S18534 | PH1318  | B58502 | F53284 | B28587     | F28587 | D28587             | A61577    | A59137             | H49048 | PA0099 | I53284 | ID            |
| hypothetical prote | cytochrome c2 - fu |        | Ig heavy chain DJ | H+-transporting tw | NADH2 dehydrogenas | urinary tract ston |        | T-cell receptor be | K allergen - ry | heavy  | Ig heavy chain J r | ptidylprolyl is | kappa-1 chain | kappa chain J | Ig kappa-1 chain J | ם      | ю.     | hain DJ | m      | 11     | 1 receptor |        | T-cell receptor be | erine pro | protein Pil - gold | epto   | 7.     |        | Description   |

RESULT 3

| <b>4</b><br>5      | 44                 | 43                 | 42                 | 41                | 40                 | 39                 | 38                 | 37                 | 36                | 35                | 34                 | 33                 | 32                 | 31                 | 30                 |
|--------------------|--------------------|--------------------|--------------------|-------------------|--------------------|--------------------|--------------------|--------------------|-------------------|-------------------|--------------------|--------------------|--------------------|--------------------|--------------------|
| 20                 | 20                 | 20                 | 20                 | 20                | 21                 | 21                 | 21                 | 21                 | 21                | 21                | 21                 | 21                 | 21                 | 21                 | 21                 |
| 20.6               | 20.6               | 20.6               | 20.6               | 20.6              | 21.6               | 21.6               | 21.6               | 21.6               | 21.6              |                   | 21.6               | 21.6               | 21.6               | 21.6               | 21.6               |
| 14                 | 14                 | 13                 | 9                  | 9                 | 18                 | 18                 | 17                 | 16                 | 15                | 15                | 14                 | 14                 | 12                 | 10                 | 10                 |
| Ν                  | N                  | N                  | N                  | N                 | N                  | N                  | N                  | N                  | N                 | N                 | Ν                  | N                  | N                  | N                  | N                  |
| G44957             | S22236             | B58533             | S10784             | D48186            | S70612             | S57518             | S50901             | E53284             | S36893            | B61457            | S59495             | PL0142             | A53524             | A39745             | S71948             |
| photosystem II oxy | lipoxygenase (EC 1 | CD61 homolog - cha | enamelin i - bovin | ATPase R1 subunit | alpha-macroglobuli | T cell receptor be | chlorophyll a/b-bi | T-cell receptor be | ribosomal protein | alpha-glucosidase | formate dehydrogen | carbon-monoxide de | ubiquinol-cytochro | endo-glucosylceram | matrix metalloprot |

# ALIGNMENTS

| Query Match 27.8*; Score 27; DB 2; Length 15; Best Local Similarity 62.5*; Pred. No. 3.2e+02; Matches 5; Conservative 1; Mismatches 2; Indels 0; Gaps 0; Qy 10 FPYTAPGT 17 | RESULT 2  PA0099  phenotipic variation protein - fungus (Fusarium sporotrichioides) (fragment)  C;Species: Fusarium sporotrichioides  C;Date: 20.Feb-1995 #sequence_revision 20.Feb-1995 #text_change 09-Jul-2004  C;Accession: PA0099  R;Chow, L.P.; Fukaya, N.; Sugiura, Y.; Ueno, Y.; Tabuchi, K.; Tsugita, A.  submitted to JIPID, October 1994  A;Description: Two dimensional polyacrylamide gel electropheresis of Fusarium sporc  A;Reference number: PA0051  A;Accession: PA0099  A;Molecule type: protein  A;Residues: 1-15 < CHO  A;Cross-references: UNIPROT:Q7M4Y7; UNIPARC:UPI000017B405 | Qy 12 YTAPGTK 18<br> | Query Match 28.9%; Score 28; DB 2; Length 15;<br>Best Local Similarity 71.4%; Pred. No. 2.1e+02;<br>Matches 5; Conservative 0; Mismatches 2; Indels 0; Gaps 0; | A;Status: preliminary A;Molecule type: DNA A;Residues: 1-15 <4RR> A;Cross-references: UNIPARC:UPI0000115418; GB:S60737; NID:g233916; PIDN:AAB19525.1; A;Note: sequence extracted from NCBI backbone (NCBIN:60737, NCBIP:60747) C;Keywords: T-cell receptor | Alexander, C.B.; Mage, R.G.<br>81-888, 1991<br>rily conserved organization and sequences of germline diversity<br>AS3284; MUID:91342695; PMID:1678859 | RESULT 1<br>153284<br>T-cell receptor beta 2 chain J region, Jbeta2.7 - rabbit<br>C;Species: Oryctolagus cuniculus (domestic rabbit)<br>C;Date: O2-May-1994 #sequence_revision 18-Nov-1994 #text_change O5-Nov-1999<br>C;Accession: I53284 |
|--|--|----------------------|--|--|---|--|
|  | sporotric  |                      |  | 5.1; PID   | ity and   |  |

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H49048
T-cell receptor beta chain V region (CDR3 junction, clone Vbeta20:1) - human (fragment C;Species: Homo sapiens (man)
C;Date: 21-Jan-1994 #sequence_revision 18-Nov-1994 #text_change 30-May-1997
C;Accession: H49048
R;Sioud, M; Kjeldsen-Kragh, J; Suleyman, S; Vinje, O.; Natvig, J.B.; Forre, O.
Eur. J. Immunol. 22, 2413-2418, 1992
A;Title: Limited heterogeneity of T cell receptor variable region gene usage in juvent, Reference number: A49048; MUID:92387250; PMID:1387614
A;Accession: H49048
A;Status: preliminary; not compared with conceptual translation
A;Cross-references: UNIPARC:UPI0000176DF0
A;Cross-references: UNIPARC:UPI0000176DF0
A;Status: sequence extracted from NCB backbone (NCBIP:113270)
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein Pil - golden needle mushroom (fragment)
C;Species: Flammulina velutipes (golden needle mushroom)
C;Date: 12-Nov-1999 #sequence_revision 12-Nov-1999 #text
C;Accession: A59137
R;Sakamoto, Y.: Ando a "
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A61577
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A; Mesidues: 1-18 < SAK>
A; Cross-references: UNIPROT:Q7M4W6; UNIPARC:UPI000017CB27
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                                                                                                                                                                                  R;Sinha, U.; Wolz, S.A.; Lad, P.J.
Int. J. Biochem. 23, 979-984, 1991
A;Title: Two new extracellular serine proteases from Streptomyces
A;Reference number: A61577; MUID:92155439; PMID:1786859
                                                                                                                                                                                                                                                                           24k serine proteinase (EC 3.4.21.-) - Streptomyces fradiae (fragment) C;Species: Streptomyces fradiae (C;Species: 28-Oct-1994 #sequence_revision 28-Oct-1994 #text_change 09-Ju
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Sakamoto, Y.; Ando, A.; Tamai, Y.; Miura, K.
ubmitted to the Protein Sequence Database, November 1999
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                                                                                       Cross-references: UNIPROT:Q7M198; UNIPARC:UP1000017AE13 Keywords: extracellular protein; hydrolase; serine prot
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Similarity 41.7%;
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62.5%;
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Pred. No. 5.7e+02;
2; Mismatches 5;
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Pred. No. 3.9e+02;
1; Mismatches 0;
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Pred. No. 3.6e+02;
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R;Toyonaga, B.; Yoshikai, Y.; Vadasz, V.; Chin, B.; Mak, T.W. Proc. Natl. Acad. Sci. U.S.A. 82, 8624-8628, 1985.
A;Title: Organization and sequences of the diversity, joining, A;Reference number: A94081; MUID:86094276; PMID:3866244
A;Accession: D28587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Toyonaga, B.; Yoshikai, Y.; Vadasz, V.; Chin, B.; Mak, T.W. Proc. Natl. Acad. Sci. U.S.A. 82, 8624-8628, 1995
A;Title: Organization and sequences of the diversity, joining, A;Reference number: A94081; MUID:86094276; PMID:3866244
A;Accession: F28587.
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                                                           A;Cross-references: UNIPARC:UPI000002FD06; GB:M14159; NID:g338852; C;Keywords: T-cell receptor
                                                                                                                                                         R;Toyonaga, B.; Yoshikai, Y.; Vadasz, V.; Chin, B.; Mak, T.W. Proc. Natl. Acad. Sci. U.S.A. 82, 8624-8628, 1985
A;Title: Organization and sequences of the diversity, joining A;Reference number: A94081; MUID:86094276; PMID:3866244
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C;Date: 16-Aug-1988 #sequence_revision 16-Aug-1988
C;Accession: B28587
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A; Residues: 1-15 < TOY>
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A; Residues: 1-16 < TOY>
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A;Residues: 1-15 <TOY>
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Similarity 57.1%;
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Ig heavy chain DJ region (clone C527-121) - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1318
R;Wasserman, R; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
J. Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymph A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Moclession: PH1318
A;Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mol. Immunol. 28, 881-888, 1991
A;Title: Evolutionarily conserved organization and sequences A;Reference number: A53284; MUID:91342695; PMID:1678859
A;Accession: F53284
A;Status: preliminary
                                                                                                                                                                                                      RESULT 11
PH1318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      36K kidney stone protein - unidentified bacterium (fragment) C;Species: unidentified bacterium C;Date: 07-Feb-1997 #sequence_revision 07-Feb-1997 #text_change 10-Jul-1998 C;Accession: B58502 R;Binette, J.P.; Binette, M.B. submitted to the Protein Sequence Database, October 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPARC:UPI000017A8D0

A;Experimental source: human kidney stone containing Ca ox.mono& dihyd, 1% struvite, Car A;Note: tentative identification of 8-Tyr and 9-Thr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 9
F53284
T-cell receptor beta 2 chain J region, Jbeta2.3 - rabbit C;Species: Oryctolagus cuniculus (domestic rabbit) C;Date: O2-May-1994 #sequence_revision 18-Nov-1994 #text. C;Accession: F53284
R;Harindranath, N.; Alexander, C.B.; Mage, R.G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-16 <HAR>
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Pred. No. 9.5e+02;
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Pred. No. 7.46
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Gene 102, 261-264, 1991
A;Title: Immunoglobulin heavy-chain joining genes in the
A:Reference number: JH0666; MUID:91340162; PMID:1908401
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S18534
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C;Superfamily: immunoglobulin V region; imm
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                       R;Brueggemann, M.; Free, J.; Diamond, A.; Howard, J.; Cobbold, Proc. Natl. Acad. Sci. U.S.A. 83, 6075-6079, 1986
A;Title: Immunoglobulin heavy chain locus of the rat: striking A;Reference number: A25941; MUID:86287397; PMID:3016742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ig heavy chain J-H1 region - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 05-Jun-1988 #sequence_revision 05-Jun-1988 #text_change 23-Jul-1999
C;Accession: A25941; JH0666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Accession: S18534
R;Haydock, S.F.; Dowson, J.A.; Dhillon, N.; Roberts, G.A.; Cortes, J.; Leadlay, P.F. Mol. Gen. Genet. 230, 120-128, 1991
A;Title: Cloning and sequence analysis of genes involved in erythromycin biosynthesis
                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-18 < LAN>
                                                                                                                                                                                                              A; Reference number: JH0666; A; Accession: JH0666
                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPARC:UPI0000114C27; dB:M13798; NID:g204707; PIDN:AAA41371.1; R;Lang, P.; Mocikat, R.
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C;Species: Saccharopolyspora erythraea
C;Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text_change 07-Apr-1994
                                                                                                                                                                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-18 < BRU>
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A;Molecule type: DNA
A;Residues: 1-17 <HAY>
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A;Reference number: S18530; MUID:92079886; PMID:1840640
A;Accession: S18534
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Ig kappa-1 chain J3 region - rabbit C;Species: Oryctolagus cuniculus (domestic rabbit) C;Species: Oryctolagus cuniculus (domestic rabbit) C;C;Accession: C20907 #sccession: C20907 R;Emorine, L.; Max, E.E. Nucleic Acids Res. 11, 8877-8890, 1983 Nucleic Acids Res. 11, 8877-8890, 1983 A;Title: Structural analysis of a rabbit immunoglobulin kappa2 J-C locus reveals multipl A;Reference number: A20907; MUID:84169523; PMID:6324107 A;Accession: C20907
                                                                                                                                                                                                                                                                          Ig kappa chain J segment (J-kappa-3) - human
C;Species: Homo sapiens (man)
C;Accession: S23640
R;Huber, C:, Klobeck, H.G.; Zachau, H.G.
Eur. J. Immunol. 22, 1561-1565, 1992
A;Title: Ongoing V(kappa) - V(kappa) recombination after formation of a productive V(kappa A;Reference number: S23637; MUID:92289816; PMID:1601042
A;Recession: S23640
A;Recession: S23640
A;Statua: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-13 :HUB>
A;Rote: the nucleotide sequence was submitted to the EMBL Data Library, November 1991
C;Keywords: heterotetramer; immunoglobulin
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S23640
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A;Molecule type: DNA
A;Residues: 1-12 <EMO>
A;Cross-references: UNIPARC:UPI000017C5CF
C;Keywords: heterotetramer; immunoglobulin
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

| 31                 | 30       | 29     | 28           | 27                | 26          | 25           | 24           | 23           | 22           | 21   | 20                 | 19           | 18           | 17           | 16   | 15                 | 14                | 13                | 12           | 11           | 10                 | 9            | 8             | 7            | σ            | σ            | 4.           | w        | N                 |                | No.         | Result |
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| Q3ZEY3_9CHAR       | ZEXI     | 3ZEW8  | Q3ZEW2_9CHAR | Q3ZEU7_9CHAR      | O89560_HHV8 | Q8L1Y8_9CYAN | Q7M263_SECCE | Q85HM7_MOTFL | Q5D4R7_9CYAN |      | CWP27_TOBAC        | Q9ZIB1_CLODI | Q5ZEY7_HUMAN | Q65YZ6_HORSE | <    | 069142_STRPY       | 714T              | 9UR63             | Q7M198_STRFR | Q7S007_NEUCR | CWP16_LYCES        | Q3TW38_MOUSE | Q7M4W6_FLAVE  | Q3ZEY6_9CHAR | Q7M4Y7_FUSSP | Q8L1Y7 PLEBO | Q3ZEV4_9CHAR | 9TWL4    | Q70Y88_9LAMI      | O46473_FELCA   | ID          |        |
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Q9TWL4;
Q1-MAY-2000, integrated into UniProtKB/TrEMBL.
Q1-MAY-2000, sequence version 1.
Q1-FEB-2006, entry version 11.
Q2-FEB-2006, entry version 11.
Q5-KDa chymotrypsin-like enzyme (Fragment).
Lucilia cuprina (Greenbottle fly) (Australian sheep blowfly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Oes
Calliphoridae; Lucilia.
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MEDLINE=95219141; |
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Nucleic Acids Res. 30:2025-2030(2002).
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MEDLINE=21970088; PubMed=11972342; DOI=10.1093/nar/30.9
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RESULT 8
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27-SEP-2005,
07-FEB-2006,
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15-DEC-2003, sequence version 1.
07-FEB-2006, entry version 6.
Phenotipic variation protein (Fragment).
Pusarium sporotrichioides.
Eukaryota; Pungi; Ascomycota; Pezizomycotina; Sordariomycetes;
Eukaryota; Pungi; Ascomycota; mitosporic Hypocreales; Fusari
                                                                                                                                                                                                                                                   NUCLECTIDE SEQUENCE.

Pereira S.L., Baker A.J.;

"Multiple gene evidence for parallel evolution and retention o ancestral morphological states in the shanks (Charadriiformes)
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                                                                                                                                           SEQÜENCE
                                                                                                                                                              Mitochondrion.
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                                                                                                                                                                                                         Distributed
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                                                                                                                                                                                                                                      Condor 107:514-526(2005).
                                                                                                                                                                                                                                                  Scolopacidae)
                                                                                                                                                                                                                                                                                                                  Archosauria; Aves;
NCBI_TaxID=161739;
                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                 Mitochondrion.
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                                                                                                                                                                                                                                                                                                                                                                                ATP synthase
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Q3ZEY6;
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RESULT 9
Q3TW38 MOUSE
ID Q3TW38 MOUSE
AC Q3TW38;
RA Carninci P., Kasukawa T., Katayama S., Gough J., Frith M.C., Maeda N., RA Oyama R., Ravasi T., Lenhard B., Wells C., Kodzius R., Shimokawa K., RA Oyama R., Ravasi T., Lenhard B., Wells C., Kodzius R., Shimokawa K., RA Davis M.C., Wilming L.G., Aidinis V., Allen J.E., Ra Davis M.J., Wilming L.G., Aidinis V., Allen J.E., Bansal M., Wilming L.G., Aidinis V., Allen J.E., Bono H., Chalk A.M., Ra Ambesi-Impiombato A., Apweiler R., Aturaliya R.N., Bailey T.L., Ra Bansal M., Choudhary V., Christoffels A., Clutterbuck D.R., Chalk A.M., Ra Crowe M.L., Dalla E., Dalrymple B.P., de Bono B., Della Gatta G., Ra Crowe M.L., Dalla E., Dalrymple B.P., de Bono B., Della Gatta G., Ra Crowe M.L., Down T., Engstrom P., Fagiolini M., Faulkner G., Ra Ai Bernardo D., Down T., Engstrom P., Fagiolini M., Faulkner G., Ra Ai Georgii-Hemming P., Gingeras T.R., Gojobori T., Green R.E., Georgii-Hemming P., Gingeras T.R., Gojobori T., Green R.E., Gustincich S., Harbers M., Hayashi Y., Hensch T.K., Hirokawa T., Ra Gustincich S., Harbers M., Hayashi Y., Hensch T.K., Jishikawa T., Ra Hill D., Huminiecki L., Iacono M., Ikoo K., Iwama A., Ishikawa T., Jakt M., Kanapin A., Katch M., Kawasawa Y., Kelso J., Kitamura H., Kaka M., Katano H., Kollias G., Krishnan S.P., Kruger A., Kummerfeld S.K.,
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Best Local S
Matches 4
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Osteoclast-like cell colon, RIKEN full-length enriched library, clone:1420034D23 product:RNA binding motif, single stranded interacting protein 1, full insert sequence. (Fragment).
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SEQUENCE
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15-DEC-2003,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                                               PubMed=16141072; DOI=10.1126/science.1112014;
                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9; Carninci P., Hayashizaki Y.; "High-efficiency full-length cDNA cloning."; Methods Enzymol. 303:19-44(1999).
                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE
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11-OCT-2005,
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Eukaryota; Fungi; Basidiomycota; Hymenomycetes; Homobasidiomycetes;
Posricales: Tricholomataceae; Flammulina.
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STRAIN=C578L/6J;
DOI=10.1126/science.1112009;
PubMed=16141073; DOI=10.1126/science.1112009;
RIKEN Genome Exploration Research Group, and Genome Science
RIKEN Genome Network Core Team) and the FANTOM Consortium;
(Genome Network Core Team) in the Mammalian Transcriptome.";
"Antisense Transcription in the Mammalian Transcriptome.";
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MEDLINE-21085660; PubMed-11217851; DOI-10.1038/35055500; Kawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito
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"Analysis of the mouse transcriptome 60,770 full-length cDNAs.";
Nature 420:563-573(2002).
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RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
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RA Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
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STRAIN=C57BL/6J;
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Distributed under the Creative Commons Attribution-NoDerivs License
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GO:0003690; F:double-stranded DNA binding;
GO:0003723; F:RNA binding; RCA.
GO:0003697; F:single-stranded DNA binding;
GO:0006260; P:DNA replication; RCA.
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A Galagan J.E., Calvo S.E., Borkovich K.A., Selker E.U., Read N.D., Jaffe D., FitzHugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B., Elkins T., Engels R., Wang S., Nielsen C.B., Butler J., Endrizzi M., Qui D., Ianakiev P., Bell-Pedersen D., Nelson M.A., Braun E.L., Zelter A., Schulte U., Kothe G.O., Jedd G., Mewes H.-W., Staben C., Marcotte E., Greenberg D., Roy A., Foley K., Naylor J., Ramorotte E., Greenberg D., Roy A., Foley K., Naylor J., Ramorotte E., Belke C., Rudd S., Frishnan D., Krystofova S., Marceli E., Blelke C., Rudd S., Frishnan D., Krystofova S., Rasmussen C., Metzenberg R.L., Perkins D.D., Kroken S., Cogoni C., Macino G., Catcheside D.E.A., Li W., Pratt R.J., Osmani S.A., Macino G., Catcheside D.E.A., Li W., Pratt R.J., Osmani S.A., Ander C., Plamann M., Seiler S., Dunlap J.C., Radford A., Aramayo R., Varden O., Plamann M., Seiler S., Dunlap J.C., Radford A., Aramayo R., Paulsen I., Sachs M.S., Lander E.S., Nusbaum C., Birren B.W.;

"The genome sequence of the filamentous fungus Neurospora crassa.";
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STRAIN=74-OR23-1A / FGSC 987;
MEDLINE=22598136; PubMed=12712197; DOI=10.1038/nature01554;
Galagan J.E., Calvo S.E., Borkovich K.A., Selker E.U., Read
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MEDLINE=97332671; PubMed=9188482; DOI=10.1074/jbc.272.25.15841;
Robertson D., Mitchell G.P., Gilroy J.S., Gerrish C., Bolwell G.
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07-FEB-2006, entry version 4.
40 kDa cell wall protein (Fragment).
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40 kDa cell wall protein.

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1705 MW;
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Pred. No. 5.5e
1; Mismatches
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s; Sordariaceae;
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Neurospora.
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RESULT 13
Q9UR63 EMENI
Q9UR63;
ID Q9UR63;
DT 01-MAY-2000, integrated into Uni
DT 01-MAY-2000, sequence version 1.
DT 07-FEB-2006, entry version 11.
DE Beta-D-FRUCTOFURANOSIDE FRUCTOHY
DE isoform (EC 3.2.1.26) (Fragment)
OS Emericella nidulans (Aspergillus)
OC Eukaryota; Fungi; Ascomycota; Pe
OC Eurotiales; Trichocomaceae; Emer
OX NCBI TaxID=162425;
RY MEDLINE=96409246; PubMed=8814228
RA Chen J.S., Saxton J., Hemming F.
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Eukaryota; Fungi; Ascomycota; Pezizomycotii
Eurotiales; Trichocomaceae; Emericella.
NCBI TaxID=162425;
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15-DEC-2003,
07-FEB-2006,
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=92155439; PubMed=1786859; DOI=10.1016/0020-711X(91)90133-8; Sinha U., Wolz S.A., Lad P.J.; Sinha U., Wolz S.A., Lad P.J.; "Two new extracellular servine proteases from Streptomyces fradiae." Int. J. Biochem. 23:979-984(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales; Streptomycineae; Streptomycetaceae; Streptomyces.
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Q7M198;
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NCE 15 AA;
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    PubMed=8814228; DOI=10.1016/0167-4838(96)00073-8;
J., Hemming F.W., Peberdy J.F.;
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1832 MW; 012D0:
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41.7%;
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66.7%;
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Pred. No. 5.5e
1; Mismatches
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012D0180A8C7089D
                                                                                                                                                 Pezizomycotina;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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No. 6.7e+03;
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5.5e+03;
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                                                                                                                                                                                                                   high molecular weight
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RESULT 14
Q714T5_9CRYP
ID Q714T5_9CRYP
AC Q714T5;
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069142_STRPY
ID 069142_STRPY PRELIMINARY; PRT; 15 AA.
AC 069142;
DT 01-AUG-1998, integrated into UniProtKB/TrEMBL.
DT 01-AUG-1998, sequence version 1.
DT 07-FEB-2006, entry version 14.
DE Protein SIC (Fragment).
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STRAIN=(CMP1319;

PubMed=14681369; DOI=10.1093/nar/gkh102;

Gueneau de Novoa P., Williams K.P.;

"The tmRNA website: reductive evolution of tmRNA in plastids and other
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Biochim. Biophys. Acta 1296:207-218(1996).
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Nucleic Acids Res. 32:D104-D108(2004).
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                                                 Streptococcus pyogenes
Bacteria; Firmicutes;
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                   NCBI_TaxID=1314;
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UENCE 15 AA; 1588 MW; 2C992B42211366BB CRC64;
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                                                                                                                  EMBL; AF064540; AAC38769.1; -; Genomic_DNA.
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                                                                                                     SEQUENCE
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MEDLINE=98298075; PubMed=9632622;
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Search completed: May 30, 2006, 15:07:40 Job time: 70.6667 secs

GenCore version 5.1.8
Copyright (c) 1993 - 2006 Biocceleration Ltd.

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Adw78189 Human met
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ALIGNMENTS

#### Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody. Canis familiaris. Dog IgE epitope recognised by monoclonal antibody 3.76, SEQ ID 9 ADR10609; ADR10609 standard; peptide; 18 15-JAN-2004; 2004WO-US003566 05-AUG-2004. WO2004065936-A2 21-OCT-2004 (first entry) ₽

The present invention relates to a novel monoclonal antibody (I) that specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. (I) is useful for testing an allergen reactivity of an IgE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut and corn allergens. The sample is a biological sample collected from a dog, cat or horse. (I) is also useful for detecting mammalian IgE and fo treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal antibodies recognise epitopes on canine IgE corresponding to amino acid residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from

for

Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian IgE or treating asthma or anaphylactic shock.

WPI; 2004-593545/57.

Hammerberg B;

16-JAN-2003; 2003US-0440472P.

(UYNC-) UNIV NORTH CAROLINA STATE

Example 6; Page 9; 14pp; English.

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ADRIULT 2
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                                                                                                                                       The present invention relates to a novel monoclonal antibody (I) that CC specifically binds to a mammalian IgE spitope, where the epitope is CC between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. CC (I) is useful for testing an allergen reactivity of an IgE sample. The CC allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut CC and corn allergens. The sample is a biological sample collected from a CC dog, cat or horse. (I) is also useful for detecting mammalian IgE and for CC treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal CC antibodies recognise epitopes on canine IgE corresponding to amino acid CC residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the CC canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from CC cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and CC of IgE from cat and horse, but did not exhibit cross-reactivity with CC either pig or human epsilon-chains of IgE. The present sequence is the CC cat IgE 3.76 recognition site.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and 3.76 were observed to have good cross-reactivity with the epsilon-chain of IgE from cat and horse, but did not exhibit cross-reactivity with either pig or human epsilon-chains of IgE.
                                                                                        Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 6; Page 9; 14pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cat IgE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2004-593545/57
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18; Conserv
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   89.7%;
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Pred. No. 6.5e-09;
   Score
Pred.
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   . 87;
   DB 8; Lo
3.1e-07;
                            Length 18;
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RESULT 3
ADR10612
ID ADR1
                                                                                                                          The present invention relates to a novel monoclonal antibody (I) that CC specifically binds to a mammalian IgE spitope, where the epitope is CC between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. CC (I) is useful for testing an allergen reactivity of an IgE sample. The CC allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut CC and corn allergens. The sample is a biological sample collected from a CC dog, cat or horse. (I) is also useful for detecting mammalian IgE and for CC treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal CC antibodies recognise epitopes on canine IgE corresponding to amino acid CC residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the CC canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from CC 3.76 were observed to have good cross-reactivity with the epsilon-chain CC of IgE from cat and horse, but did not exhibit cross-reactivity with CCC sheep IgE 3.76 recognition site.
                                                             Matches
                                                                        Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian I or treating asthma or anaphylactic shock.
                                                                                                                        Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; Page 9; 14pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sheep IgE epitope recognised by monoclonal antibody 3.76, SEQ ID 12.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hammerberg
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                                                                            Similarity
                            VDGQKATNIFPYTAP 15
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VDGQEDRNLFSYTAP
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                                                             Conservative
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                                                                        56.7%;
66.7%;
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                                                           Score 55; DB 8;
Pred. No. 0.071;
2; Mismatches
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                                                                                          DB 8;
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RESULT 4 ADC56890

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ADC64569
ID ADC64
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DT 18-C
DT 18-C
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                                                                                                                                                                                                                                                                                                                                     Matches
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                      08-NOV-2001; 2001US-00052788
                                            08-NOV-2001; 2001US-00052788
                                                                    08-MAY-2003
                                                                                          US2003087314-A1
                                                                                                                                    Horse; immunoglobulin E; IgE; heavy chain; immunogen; allergy
                                                                                                                                                           Horse immunoglobulin E, IgE, heavy chain immunogenic peptide P5
                                                                                                                                                                                   18-DEC-2003
                                                                                                                                                                                                          ADC64569;
                                                                                                                                                                                                                              ADC64569 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 6; SEQ ID NO 7; 33pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New human macroprotein-36.41, (recombinant production, useful and digestive ulcers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-269480/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mao Y,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human; protein 36-41; arrhythmia; asthma; dementia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptide fragment Seq
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 (REGC ) UNIV CALIFORNIA
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                                                                                                                                                                                                                                                                                                               1 VDGQKATNIFPYTAP 15
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                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                38.1%;
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                                                                                                                                                                                                                                                                                                                                                 Score 37;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoding polynucleotide,
l for treating dementia, a
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                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                            B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                human protein 36-41, and the
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                                                                                                                                                                                                                                                                                                                                                            Length 15
                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , antagonist
arrhythmia,
                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                     Gaps
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asthma
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Length 15

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cc antibody that specifically binds to equine immunoglobulin (Ig)E), the six cc polypeptides are not explicitly identified in the specification. Also cc included are a composition comprising an antibody that specifically binds to equine IgE made by the process of immunising an animal with a polypeptide at least 80% identical to (S1)-(S6), an antibody that specifically binds to equine IgE made by the process of immunising an cc antibody that specifically binds to equine IgE (involving immunising an cc amimal with a composition further comprising an isolated polypeptide (the camino acid sequence of the polypeptide is at least 80% identical to (S1)-(S6), making an cc amino acid sequence of the polypeptide is at least 80% identical to (S1)-(C (S6)), and collecting antiserum from the animal) and a kit for detection of equine IgE in a biological sample comprising the antibody and means (C thus forming an antigen/antibody complex, and detecting the polypeptide (serum) which involves contacting the sample with the antibody, to equine IgE protein in a biological sample (serum) which involves contacting the sample with the antibody, to equine IgE and a server as a diagnostic test for altered such that the complex can be detected. The complex. The antibody and antigen are complex can be detected. The complex is detected using a second labelled can be an erver as a diagnostic test for allergy. The present complex is a Horse immunoglobulin E, IgE, heavy chain immunogenic peptide from the middle portion of the C2 region.
Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           identical to 6 (S1-S6), 15 amino acid peptide sequences derived from equine immunoglobulin E (the composition induces production of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to an immunogenic composition comprising isolated polypeptide having an amino acid sequence that is at identical to 6 (S1-S6), 15 amino acid peptide sequences derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 8; 14pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunogenic composition c polypeptide that induces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 to equine immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-765437/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gershwin LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pettigrew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprising an isolated equine immunoglobulin E production of antibodies which specifically bind 1 E.
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RESULT 6
ADR10611
ID ADR1
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XX ADR1
XX ADR1
XX HOTB
XX HOTB
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Best Local S
Matches
                                      15-JAN-2004; 2004WO-US003566.
                                                                                                               Equus caballus.
                                                                                                                                                 Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody;
                                                                                                                                                                                      Horse IgE epitope recognised
                                                                                                                                                                                                                 21-OCT-2004
                                                                                                                                                                                                                                          ADR10611;
                                                                                                                                                                                                                                                                 ADR10611 standard; peptide; 18
              16-JAN-2003; 2003US-0440472P
                                                                                      WO2004065936-A2
                                                                                                                                                                                                                                                                                                                                                                                Local Similarity hes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                        1 VDGQKATNIFP 11
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                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                            35.1%;
54.5%;
                                                                                                                                                                                                                                                                                                                                                                                            Score 34; DB 7;
Pred. No. 1.9e+02;
                                                                                                                                                                                      by monoclonal antibody 3.76, SEQ ID 11.
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Best Local
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                                                                                                                                                                                                                               DEF domain; MAP kinase; cytostatic; cardiovascular-gen.; cardiant; vasotropic; hypotensive; antiarteriosclerotic; antianflammatory; antiallergic; immunosuppressive; antibacterial; antianflammatory; dermatological; antidiabetic; gastrointestinal-gen.; antialcer; thrombolytic; neuroprotective; ophthalmological; antirheumatic; antipyretic; uropathic; antipsoriatic; hepatotropic; antianemic; muscular-gen.; thycomimetic; antithyroid; gynecological; nephrotropic; hepatotropic; virucide, anti-HIV; anabolic; hypertensive; anorectic; endocrine-gen.; neuroleptic; nootropic; antiparkinsonian; anticonvulsant; antidepressant; antiaddictive; sedative; hypotic; CNS-gen.; antipiertility; cancer; cardiovascular disease; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a novel monoclonal antibody (I) that specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 336-374 of mammalian IgE, e.g. dog IgE. (I) is useful for testing an allergen reactivity of an IgE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut and corn allergens. The sample is a biological sample collected from a dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal antibodies recognise epitopes on canine IgE corresponding to amino acid residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and cat, horse, pig and human by 5.91 and 3.76 was examined.
                      03-JUL-2003; 2003US-0484761P
                                                          02-JUL-2004; 2004WO-US021514
                                                                                                                                      WO2005007090-A2
                                                                                                                                                                             Homo
                                                                                                                                                                                                                 metabolic disorder; neuropathy; sleep disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human metabolic therapy target peptide PAGT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-APR-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADW78189;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADW78189 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 18 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.76 were observed to have good cross-reactivity with the epsilon-chain of IgE from cat and horse, but did not exhibit cross-reactivity with either pig or human epsilon-chains of IgE. The present sequence is the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 6; Page 9; 14pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian I or treating asthma or anaphylactic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hammerberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               horse IgE
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                                                                                                                                                                           sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 VDGQKATNIFP 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IDGOKVDEQFP 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.76 recognition site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 35.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 34;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 8;
2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IgE
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RESULT 8
AAU7650
ID AAU7
XX AAU7
XX AAU7
XX AAU7
XX BAT1
XX BA
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Best Local S
Matches 8
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                                                                                                                                                                                                                  multiple sclerosis; i
neurological disorder
                                                                                                                                                                                                                                                                                        complementarity determining region; neuroprotective; antitipsoriatic;
immunostimulant; cytostatic; anti-microbial; psoriasis; infection;
                                                                                                                                                                                                                                                                                                                                                              Human; antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Anti-Interleukin-12 (IL-12) antibody CDR3 light chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAU76520 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 14; Page 64; 104pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the compound.
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                              anti-interleukin-12; CDR; light chain;
                                                                                                                                                                                                                                                       immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    33.5%;
                                                                                                                                                                                                                                                       disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 32.5; DB 9;
Pred. No. 3.4e+02;
D; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                  circulatory;
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WO200212500-A2 Homo sapiens.

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Best Local S
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29-SEP-2000;
01-AUG-2001;
                                                                                                                                                                                                                         Human; sialoadhesin factor-3; SAF-3; therapy; cancer; inflammation; autoimmunity; allergy; asthma; infection; central nervous system; CNS; rheumatoid arthritis; multiple sclerosis; stem cell mobilisation; AIDS; haematopoietic development; anaemia; chemoprotective agent; cytostatic; immunoglobulin; complementarity determining region; CDR; protozoacide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           determining region (CDR) of a heavy or light chain, a heavy chain or light chain variable region, or a heavy chain or light chain constant region. The anti-IL-12 antibodies are useful in methods for diagnosing or treating IL-12 related conditions, e.g. psoriasis or multiple sclerosis. The antibodies are also useful for treating immune, cardiovascular, infectious, malignant or neurological disorders or diseases. The present sequence represents the amino acid sequence of human anti-Interleukin- 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New mammalian anti-IL-12 antibodies, useful for diagnosing or treating IL-12 related conditions, e.g. psoriasis or multiple sclerosis, as well as other for treating immune, infectious, malignant or neurological
                                         24-MAY-2000;
                                                                    24-MAY-2001; 2001WO-US016864.
                                                                                                 29-NOV-2001
                                                                                                                            WO200190193-A1
                                                                                                                                                      Homo sapiens
                                                                                                                                                                                    monoclonal antibody
                                                                                                                                                                                             antiinflammatory; immunosuppressive; anti-HIV; fungicide; neuroprotective; light chain variab
                                                                                                                                                                                                                                                                                                                Human mAb
                                                                                                                                                                                                                                                                                                                                           26-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                AAE15818 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 41; Page 93; 96pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Giles-Komar J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-AUG-2001; 2001WO-US024720.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CENZ ) CENTOCOR INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention relates to novel isolated mammalian anti-interleukin-12 antibodies. The antibodies comprise at least one complementarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
5; Conserv
SMITHKLINE SMITHKLINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antibody CDR3 light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LYGYIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NIFPYT 13
                                                                                                                                                                                                                                                                                                             12B1 VK complementarity determining region (CDR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000US-0223358P
2000US-0236827P
2001US-00920262
                                         2000US-00577930.
                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9
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BEECHAM CORP
BEECHAM PLC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           33.0%;
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3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 32; DB Pred. No. 2.1e 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peritt D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.1e+06; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Scallon B,
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••
                                                                                                                                                                                                             antibacterial; virucide;
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RESULT 10
AEF76312
ID AEF76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                         08-JUL-2005; 2005WO-US024260
                                                                                                                                                                                                                                                                                                                                           antibody; prostate tumor; cytostatic; andrology; genitourinary disease; neoplasm; antibody identification; antibody therapy; light chain.
                                                                                                                                                                                                                                                                                                                                                                                                   Prostate cancer cell antibody light chain CDR3 SEQ ID NO:65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to monoclonal antibodies that bind sialoadhesin factor-3 (SAF-3). SAF-3 antibody is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 13; Page 67; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treating or preventing cancer, inflammation, autoimmunity, allergy, asthma, rheumatoid arthritis, multiple sclerosis, AIDS and infections.
                                   WPI; 2006-173528/18.
                                                                                                                                                                                                                                                                   WO2006017173-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                          20-APR-2006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AEF76312 standard;
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                                                                                                                                                   10-JUL-2004; 2004US-0586811P
                                                                                                             (ALEX-) ALEXION PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               monoclonal antibody that binds to human sialoadhesin factor-3 for
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5; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                       Xin H,
                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                33.0%;
                                                                           Yantiri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 32;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                           Siva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . 2.1e+06;
ches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       to human
for treating or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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0,

New antibody that binds to a prostate cancer cell, useful for detecting

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RESULT 11
ADT40399
ID ADT404
XX ADT40
XX ADT40
XX ADT40
XX YITUG
XX VITUG
XX VITUG
XX SARS
XX WO200
XX SARS
XX WO200
XX SARS
PN WO201
XX O7-OX
XX O
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Best Local S
Matches 5
Chan K, G
                                                                                                                                                                                                                                                                                                                                                                              24-MAR-2003;
26-MAR-2003;
02-APR-2003;
03-APR-2003;
                                                                                                                                                                                                                                                                                                                                            08-APR-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Virucide; Severe Acute Respiratory Syndrome; SARS; vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hSARS virus peptide, SEQ ID 1387.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-DEC-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9
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                                                                                                                                                                                                                                                                    23-APR-2003;
                                                                                                                        (UYHK-) UNIV HONG
                                                                                                                                                                                                                                                                                                      14-APR-2003;
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2003US-0459931P.
2003US-0459931P.
2003US-0460357P.
2003US-0461265P.
2003US-0461265P.
2003US-0464886P.
2003US-0464886P.
2003US-0465738P.
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                                                                                                                        KONG.
                                                    Nicholls JM,
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Pred. No. 2.1e
1; Mismatches
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                                                    Peiris JSM,
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2.1e+06;
ches 0; Indels
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                                                    Yuen
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The present invention relates to novel human Severe Acute Respiratory Syndrome (hSARS) viral nucleic acid and protein sequences derived from a Chars virus having China Center for Type Culture Collection Deposit CC Accession No. CCTCC-V200303. The present invention also relates to novel nucleic acid molecules (1; ADT41483 or ADT41485) encoding a nucleocapsid-CC (N) or spike (S) gene protein of a hSARS virus. Also disclosed are methods for detecting the presence of a N- or S-gene of the hSARS virus or of the protein in a biological sample and identifying a subject confected with the hSARS virus. The hSARS virus, nucleic acid and protein sequences are useful as vaccines for diagnosing or treating SARS. They care also useful in clinical and scientific research applications. The hSARS virus genome (ADT39027) was obtained and the amino acid sequences of all three reading frames were deduced from the complementary strand. ADT40501 is the full-length protein encoded by the first reading frame of the complementary strand and ADT40121. ADT40501 are the peptides from the complementary strand and ADT40121. ADT40501 is the full-length protein encoded by the first protein encoded by the first protein encoded by the full-length protein encoded by the third reading frame of the complementary strand and ADT40977 is the full-length protein encoded by the third reading frame of the complementary strand and ADT40977 is the full-length protein encoded by the third reading frame of the complementary strand and ADT40977.
New nucleic acid molecule encoding nucleocapsid- or spike-gene protein of a human Severe Acute Respiratory Syndrome (hSARS) virus, useful for diagnosing and treating SARS.
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Best Local S
Matches 6
        26-MAR-2003;
02-APR-2003;
03-APR-2003;
08-APR-2003;
14-APR-2003;
23-APR-2003;
05-MAY-2003;
16-MAY-2003;
                                                                                                     24-MAR-2004;
                                                                                                                                                                            virucide; vaccine; dete real-time quantitative
                                                                                                                                                                                                        SARS virus complementary DNA strand reading frame 1 protein #279
                                                                                                                                                                                                                                             ADS79816;
                                                                                                                                                                                                                                                               ADS79816 standard; protein; 11
                                                                                 24-MAR-2003;
                                                                                                                                          WO2004085455-A1
                                                                                                                                                            SARS coronavirus
                                                                                                                                                                                                                           30-DEC-2004 (first entry)
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                                                                                                                                                                                                                                                                                                             GÓSOSNILPOT 11
               2003US-0459931P.
2003US-0460357P.
2003US-0461265P.
2003US-0461865P.
2003US-0464886P.
2003US-046488139P.
                                                                                                     2004WO-CN000247.
                                                                                 2003US-0457031P
                                                                                                                                                                            detection; severe acute respiratory syndrome; tive polymerase chain reaction; SARS.
                                                                                                                                                                                                                                                                                                                                                          33.0%;
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Pred. No. 3e+C
1; Mismatches
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3e+02;
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ID ADT37
XX ADT37
AC ADT37
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XX 30-DE
XX Viruc
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XX 07-OC
XX 07-OC
XX 24-MA
PR 26-MA
PR 02-AE
PR 03-AE
PR 03-AE
PR 14-AE
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Matches 6
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26-MAR-2003;
02-APR-2003;
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14-APR-2003;
23-APR-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ameliorating, or preventing the virus causing overs syndrome in humans using a real-time quantitative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated nucleic acid molecule useful for detecting, treating, ameliorating, or preventing the virus causing severe acute respiratory
                       New isolated human severe acute respiratory syndrome useful as vaccine for diagnosing or treating SARS or scientific research applications.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-OCT-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Virucide; Severe Acute Respiratory Syndrome; SARS; vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 11
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                                                                                                                                                                       Leung
                                                                                                                                                                                                   Chan K,
                                                                                                                                                                                                                                                                                                                                                                        03-APR-2003;
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                                                                                                                                                                                                                                                                                             ; 2003US-0457730P.
; 2003US-0459931P.
; 2003US-0460357P.
; 2003US-0461265P.
; 2003US-0462805P.
; 2003US-0464886P.
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Pred. No. 3e+C
1; Mismatches
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3e+02;
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                                                   (hSARS) virus, in clinical an
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RESULT 14
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XX AAMS2
XX AAMS2
XX CCR5
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21-MAR-2000;
21-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCR5 binding peptide; CC chemokine receptor 5; human; HIV infection; human immunodeficiency virus; AIDS; acquired immunodeficiency syndrodrug screening; identification; phage display library.
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                                                                                                Claim
                                                                                                                                                   Identifying CC chemokine receptor 5 binding compound for treating AIDS, comprises binding a molecule from library to a molecule having binding property corresponding to CCR5 and identifying bound molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-MAR-2001;
20-MAR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAM52787 standard; peptide; 12
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                                                                                                                                                                                                                                                                                WPI; 2002-010610/01.
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                                                                                                                                                                                                                                                                                                                                                                                                       (CONS-)
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                                                                                                                                                                                                                                                                                                                                          JJ,
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6; Conservative
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; 2000US-0190996P.
; 2000US-0191299P.
; 2001US-00813448.
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                                                                                                                                                                                                                                                                                                                                          Wilson CJ,
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Pred. No. 3e+0
1; Mismatches
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The invention relates to a method for identifying a CC chemokine receptor 5 (CCR5). The method involves

binding compound for screening a library

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RESULT 15
ADV13048
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Matches
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The invention relates to characterizing phosphorylated polypeptides in a sample comprising digesting the polypeptides with a protease thus generating test peptides, and collecting a fraction of test peptides that enriched for positively charged peptides. Also included are a method (comprising determining the presence, absence or level of one ore more phosphorylated peptides as identified above in cells having a cell state and determining the degree of correlation between the presence, absence or level of phosphorylated polypeptide with the cell state), an isolated peptide of 5-50 amino acids comprising an amino acid sequence that is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Characterizing phosphorylated polypeptides in a sample comprises digesting the polypeptides with a protease thus generating test pand collecting a fraction of test peptides that enriched for posicharged peptides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 16; Page 79; 123pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2005-031720/03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alzheimers disease; neuroprotective; nootropic; degeneration; tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human phosphorylated peptide
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Pred. No. 3.3e+02;
2; Mismatches 1
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enriched for positively
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CC (and which comprise a phosphorylation site within the subsequence), an CC isolated polypeptide selected from any of the polypeptides listed in the CC specification and is modified at a modification site, an isolated peptide CC comprising a mass spectral peak signatures. a method for identifying a CC treatment that modulates phosphorylation of an amino acid in a target CC polypeptide, a method for generating a peptide standard, a pair of CC peptide standards comprising the peptide obtained (where the peptide is CC phosphorylated and a corresponding peptide comprising an identical amino acid sequence but which is not phosphorylated), a system (comprising a CC computer memory comprising data files storing information relating to the identifying characteristics of positively charged peptides, and a data canalysis module capable of executing instructions for organizing and/or capating to the identifying characteristics of positively charged peptides and comprising characteristics of positively charged comprising data crelating to the identifying characteristics of positively charged peptides and comprising instructions for organizing and/or searching the capable of executing instructions for organizing the comprising characteristics of positively charged competides and comprising instructions for organizing and/or searching the capable of characterizing phosphorylated polypeptides in a computer competides in a sample. The present sequence is a peptide from a human phosphoprotein, containing a phosphorylation site, identified by the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 subsequence
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á 11 PYTAPGT Similarity 6; Conserv Conservative

Matches Query Match Best Local

Sequence 12

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Score 32; I Pred. No. 3. Mismatches

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PATAPGT

Search completed: May 30, Job time: 64.6667 secs 2006, 15:04:08

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Result
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Maximum Match 100%
Listing first 45 s
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Maximum DB
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Perfect score:
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      protein search, using
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Match Length
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1: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US09_PUBCOMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

6: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US11_PUBCOMB.pep:*
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    Copyright
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US-10-052-788-5
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US-10-975-165-11
US-09-920-262A-6
US-10-975-843-6
US-10-975-740A-6
US-10-975-740A-6
US-10-975-708-6
US-10-975-708-6
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US-10-975-708-6
US-10-808-187-1387
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US-10-808-187-1387
US-10-808-185-1037
US-10-300-61-155-1
US-10-300-694A-64
US-10-300-61-1527
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  9, Appli
12, Appl
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US-10-758-165-9
      Query Match
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|            | 45                 | 44                | <b>4</b> 3        | 42                | 41                | 40                | 39                | 38                | 37                | 36              | 35                | 34                | 33                | 32                 | 31                 | 30                 | 29                | 28                |
|------------|--------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-----------------|-------------------|-------------------|-------------------|--------------------|--------------------|--------------------|-------------------|-------------------|
|            | 28                 | 28                | 28                | 28                | 28                | 29                | 29                | 29                | 29                | 29              | 29                | 29                | 29                | 29                 | 29                 | 29                 | 29                | 30                |
|            | œ                  | ന                 | 28.9              | æ                 | æ                 | S                 | io                | S                 | S                 | S               | 29.9              | ø                 | S                 | σ                  | w                  | Ø                  | 29.9              | 30.9              |
|            | 9                  | 9                 | 9                 | 9                 | 9                 | 18                | 17                | 17                | 15                | 15              | 12                | 12                | 12                | 9                  | 9                  | 9                  | 9                 | 18                |
|            | v                  | 4.                | 4                 | 4                 | 4                 | ហ                 | 4.                | w                 | ហ                 | 4               | თ                 | v                 | ហ                 | σ                  | σ                  | თ                  | ហ                 | ហ                 |
| ALIGNMENTS | US-10-482-284A-186 | US-10-160-505-34  | US-10-688-015-34  | US-10-449-379-34  | US-10-160-506-34  | US-10-846-548A-1  | US-10-306-631-76  | US-09-992-896-51  | US-10-720-831-20  | US-10-783-455-9 | US-11-171-567-39  | US-10-996-316-39  | US-10-894-672-39  | US-11-009-769A-214 | US-11-009-873A-214 | US-11-009-840A-214 | US-10-482-630-40  | US-10-979-183-360 |
|            | Sequence 186, App  | Sequence 34, Appl | Sequence 34, Appl | Sequence 34, Appl | Sequence 34, Appl | Sequence 1, Appli | Sequence 76, Appl | Sequence 51, Appl | Sequence 20, Appl | *               | Sequence 39, Appl | Sequence 39, Appl | Sequence 39, Appl | Sequence 214, App  | Sequence 214, App  | •                  | Sequence 40, Appl | Sequence 360, App |

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Sequence 10, Application US/10758165

Publication No. US20050196816A1

GENERAL INFORMATION:
APPLICANT: Hammerberg, Bruce
ITITLE OF INVENTION: INMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR FILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin version 3.2
SEQ ID NO 10
LENGTH: 18
TYPE: PRT
COCANICAL Selic Catus
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Publication No. US20050196816A1

GENERAL INFORMATION:
APPLICANT: Hammerberg, Bruce
TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN
FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR PILING DATE: 2003-01-16
NUMBER OF SED ID NOS: 16
SOFTWARE: PatentIn version 3.2
ILENGRAD: 10
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US-10-758-165-10
                                                   US-10-758-165-10
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TYPE: PRT
ORGANISM: Canis familiaris
                                                                               ORGANISM: Felis catus
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Score 87;
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  Length 18;
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DB 5;

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                                                                                                                                                                                                                     FILE REFERENCE: 3742.1000-000

CURRENT APPLICATION NUMBER: US/10/856,118

CURRENT FILING DATE: 2004-05-27

PRIOR APPLICATION NUMBER: PCT/GB02/005411

PRIOR FILING DATE: 2002-12-02

PRIOR APPLICATION NUMBER: GB0128733.3

PRIOR APPLICATION NUMBER: US 60/334,649

PRIOR APPLICATION NUMBER: US 60/334,649

PRIOR FILING DATE: 2001-11-30

PRIOR FILING DATE: 2001-11-30

PRIOR FILING DATE: 2001-11-30

NUMBER OF SEQ ID NOS: 41

SOPTWARE: FASCESQ for Windows Version 4.0

SEQ ID NO 34

LENGTH: 15
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US-10-758-165-12
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US-10-758-165-12
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                                                                                                                               ; OTHER INFORMATION: Flanking sequence US-10-856-118-34
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APPLICANT: Hammerberg, Bruce
TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR FILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 16
                                                   Query Match 38.1%;
Best Local Similarity 55.6%;
Matches 5; Conservative
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Best Local Similarity
Matches 10; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 100.0%; I Matches 16; Conservative 0;
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SEQ ID NO 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Laidlaw, Stephen
APPLICANT: Skinner, Mike
APPLICANT: Hill, Adrian V.S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Skinner, Mike
APPLICANT: Hill, Adrian V.S.
APPLICANT: Gilbert, Sarah C.
APPLICANT: Anderson, Richard
TITLE OF INVENTION: Vaccine
                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
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                 TNIFPYTAP 15
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                                                                       Score 37; DB
Pred. No. 58;
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Pred. No. 0.079;
2; Mismatches
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                                                     Mismatches
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hes 0;
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SEQ ID NO 11
LENGTH: 18
TYPE: PRT
ROBANISM: Equus caballus
US-10-758-165-11
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US-09-920-262A-6
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US-10-758-165-11
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Best Local S
Matches 6
                 Sequence 6, Application US/09920262A Publication No. US20030124123A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 11, Application US/10758165 Publication No. US20050196816A1 GENERAL INFORMATION:
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LENGTH: 15
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APPLICANT: Gershwin, Laurel J.
APPLICANT: Pettigrew, Howard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Pettigrew, Howard David
APPLICANT: Kalina, Warren V.
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Epsilon Immunoglobulin Chain Derived Peptides for
TITLE OF INVENTION: Induction of Anti-IgE Antibodies
FILE REFERENCE: 023070-121000US
CURRENT APPLICATION NUMBER: US/10/052,788
CURRENT FILING DATE: 2001-11-08
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 5, Application US/10052788 Publication No. US20030087314A1
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TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
APPLICANT: Shealy, David
                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/440,472 PRIOR FILING DATE: 2003-01-16
                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence:epitope peptide OTHER INFORMATION: P5, middle portion of C2 of equine IgE epsilon OTHER INFORMATION: heavy chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
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                                                                                                                                                                                                                           Similarity 54.8
6; Conservative
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54.5%;
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Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                 Score 34; DB 5;
Pred. No. 2.2e+0
                                                                                                                                                                                                                                 Mismatches
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APPLICANT: APPLICANT:

Knight, David Scallon, Bernie Giles-Komar, Jill

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US-10-975-883-6
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Best Local Similarity
The 5; Conserva
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US-10-912-994-6
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                                                                                 RESULT 9
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-912-994-6
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Sequence 6, Application US/10975883 Publication No. US20050112127A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver 3.1 SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 6, Application US/10912994 Publication No. US20050002937A1 GENERAL INFORMATION:
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Best Local Similarity 83.3%;
Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Knight, David
APPLICANT: Peritt, David
APPLICANT: Scallon, Bernie
APPLICANT: Shealy, David
TITLE OF INVENTION: ANTI-IL-12 ANTIBODIES,
FILE REFERENCE: CERRO24BDIV1
CURRENT APPLICATION NUMBER: US/10/912,994
CURRENT FILING DATE: 2004-08-06
                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/223,358
PRIOR FILING DATE: 2000-08-07
PRIOR APPLICATION NUMBER: US 60/236,827
PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: US 09/920,262
PRIOR APPLICATION NUMBER: US 09/920,262
PRIOR FILING DATE: 2001-08-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Peritt, David
TITLE OF INVENTION: ANTI-IL-12 ANTIBODIES, COMPOSITIONS, METHODS AND USES FILE REFERENCE: CEN0248
FILE REPERENCE: CEN0248
CURRENT APPLICATION NUMBER: US/09/920,262A
CURRENT FILING DATE: 2002-05-06
PRIOR APPLICATION NUMBER: 60/223,358
PRIOR FILING DATE: 2000-08-07
PRIOR PILING DATE: 2000-08-07
PRIOR PILING DATE: 2000-08-07
PRIOR FILING DATE: 2000-09-29
NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Ver 3.1
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TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 NIYPYT 9
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                                                                                                                                                                                                                                        33.0%;
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Pred. No. 1.9e+06;
                                                                                                                                                                                                                                        Score 32; DB 5;
Pred. No. 1.9e+06
                                                                                                                                                                                                                          Mismatches
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RESULT 11 US-10-975-708-6

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Sequence 6, Application US/10975708 Publication No. US20050214293A1 GENERAL INFORMATION:

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Query Match
Best Local Similarity
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TITLE OF INVENTION: ANTI-IL-12 ANTIBODIES AND USES
FILE REFERENCE: CEN0248DIV04
CURRENT APPLICATION NUMBER: US/10/975,883
CURRENT FILING DATE: 2004-10-28
PRIOR APPLICATION NUMBER: US 60/223,358
PRIOR TILING DATE: 2000-08-07
PRIOR APPLICATION NUMBER: US 60/236,827
PRIOR APPLICATION NUMBER: US 60/236,827
PRIOR APPLICATION NUMBER: US 99/920,262
PRIOR FILING DATE: 2000-09-29
PRIOR FILING DATE: 2001-08-01
NUMBER: OF SEQ ID NOS: 15
NUMBER: OF SEQ ID NOS: 15
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                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 09/920,262
PRIOR FILING DATE: 2001-08-01
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver 3.1
SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 6, Application US/10975740A Publication No. US20050196838A1
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SEQ ID NO 6
LENGTH: 9
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Best Local
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TITLE OF INVENTION: ANTI-IL-12 ANTIBODIES AND NUCLEIC ACIDS
FILE REFERENCE: CENO248DIV03
CURRENT APPLICATION NUMBER: US/10/975,740A
CURRENT FILING DATE: 2004-01-28
PRIOR APPLICATION NUMBER: US 60/223,358
PRIOR PILING DATE: 2000-08-07
PRIOR PILING DATE: 2000-09-29
PRIOR FILING DATE: 2000-09-29
PRIOR FILING DATE: 2000-09-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Giles-Komar, Jill
APPLICANT: Knight, David
APPLICANT: Peritt, David
APPLICANT: Scallon, Bernie
APPLICANT: Shealy, David
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                                                                                                                                                              TYPE: PRT
ORGANISM: Homo sapiens
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                               8 NIFPYT 13
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                                                                                        33.0%;
83.3%;
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Pred. No.
                                                                                        Score 32; DB 5;
Pred. No. 1.9e+06;
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CURRENT APPLICATION NUMBER: US/10/975,708
CURRENT FILING DATE: 2004-10-28
PRIOR APPLICATION NUMBER: US 60/223,358
PRIOR FILING DATE: 2000-08-07
PRIOR PRIOR PRIOR NUMBER: US 60/236,827
PRIOR FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: US 09/920,262
PRIOR APPLICATION NUMBER: US 09/920,262
PRIOR FILING DATE: 2001-08-01
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PATENTIN Ver 3.1
SEQ ID NO 6
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US-10-808-187-1387
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US-10-975-708-6
                                                   Sequence 1387, Application US/10808187 Publication No. US20050009009A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 14
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Publication No.
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Best Local Similarity
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APPLICANT: Kikly, Kristine K.
APPLICANT: Abrahmson, Julie
TITLE OF INVENTION: Sialoadhesin Factor-3 Antibodies
FILE REFERENCE: GH50019-1C1
CURRENT APPLICATION NUMBER: US/10/246,019A
CURRENT FILING DATE: 2002-09-18
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Best Local Similarity
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APPLICANT: PEIRIS, JOSEPH S. M. APPLICANT: YUEN, KWOK YUNG APPLICANT: POON, LIT MAN
                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: USSN 09/577,930
PRIOR FILING DATE: 2000-01-22
PRIOR APPLICATION NUMBER: USSN 09/046,736
PRIOR FILING DATE: 1998-03-24
PRIOR APPLICATION NUMBER: USSN 60/041,885
PRIOR FILING DATE: 1997-04-02
NUMBER OF SEQ ID NOS: 14
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APPLICANT: Scallon, Bernie
TITLE OF INVENTION: ANTI-IL-12 ANTIBODIES AND COMPOSITIONS
FILE REFERENCE: CEN0248DIV02
                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Human
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No. US20060073133A1
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Pred. No. 1.9e
1; Mismatches
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1.9e+06;
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1.9e+06;
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APPLICANT: GUN, YI
APPLICANT: CHAN, KWOK HUNG
APPLICANT: CHAN, KWOK HUNG
APPLICANT: CHAN, KWOK HUNG
APPLICANT: NICHOLLS, JOHN M.
APPLICANT: NICHOLLS, JOHN M.
TITLE OF INVENTION: A HIGH-THROUGHPUT DIAGNOSTIC ASSAY FOR THE HUMAN VIRUS
TITLE OF INVENTION: CAUSING SEVERE ACUTE RESPIRATORY SYNDROME (SARS)
TITLE OF INVENTION: CAUSING SEVERE ACUTE RESPIRATORY SYNDROME (SARS)
TILLE OF INVENTION CAUSING SEVERE ACUTE RESPIRATORY SYNDROME (SARS)
TILLE OF INVENTION NUMBER: US/10/807,807
CURRENT APPLICATION NUMBER: US/10/807,807
CURRENT FILING DATE: 2003-03-24
PRIOR APPLICATION NUMBER: 60/457,730
PRIOR FILING DATE: 2003-03-26
PRIOR FILING DATE: 2003-04-03
PRIOR APPLICATION NUMBER: 60/459,931
PRIOR APPLICATION NUMBER: 60/459,931
PRIOR FILING DATE: 2003-04-03
PRIOR FILING DATE: 2003-04-03
PRIOR FILING DATE: 2003-04-03
PRIOR FILING DATE: 2003-04-03
PRIOR FILING DATE: 2003-04-04
PRIOR FILING DATE: 2003-04-14
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LENGTH: 11
TYPE: PRT
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PRIOR FILING DATE: 2003-04-08
PRIOR APPLICATION NUMBER: 60/462,805
PRIOR FILING DATE: 2003-04-14
PRIOR APPLICATION NUMBER: 60/468,139
PRIOR FILING DATE: 2003-05-05
PRIOR APPLICATION NUMBER: 60/464,886
PRIOR APPLICATION NUMBER: 60/464,886
PRIOR APPLICATION NUMBER: 60/464,886
PRIOR APPLICATION NUMBER: 60/464,886
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Publication No. US20050181357A1
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APPLICANT: PEIRIS, JOSEPH S. M.
APPLICANT: YUEN, KWOK YUNG
APPLICANT: POON, LIT MAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/471,200 PRIOR FILING DATE: 2003-05-16 NUMBER OF SEQ ID NOS: 2476
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TITLE OF INVENTION: A DIAGNOSTIC ASSAY FOR THE HUMAN VIRUS CAUSING SEVERE ACUTE
TITLE OF INVENTION: RESPIRATORY SYNDROME (SARS)
FILE REFERENCE: V9661.0078
CURRENT APPLICATION NUMBER: US/10/808,187
CURRENT FILING DATE: 2004-03-24
PRIOR FILING DATE: 2004-03-24
PRIOR FILING DATE: 2003-03-24
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PRIOR FILING DATE: 2003-04-02
PRIOR APPLICATION NUMBER: 60/460,357
PRIOR FILING DATE: 2003-04-03
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APPLICATION NUMBER: 60/464,886
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Similarity 54.5%;
6; Conservative
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Pred. No. 2.8e+02;
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APPLICANT: Chye, Mee Lee
APPLICANT: Li, Hong Ye
APPLICANT: Li, Hong Ye
APPLICANT: Ramalingam, Sathiskumar
APPLICANT: Poon, Leo Lit Man
APPLICANT: Poon, Leo Lit Man
APPLICANT: Peiris, Joseph Sriyal Malik
TITLE OF INVENTION: Sequences and Methods of Use Thereof For Immunization Against SA
FILE REPERENCE: 2587/73166/RDK
CURRENT FILING DATE: 2004-12-03
PRIOR APPLICATION NUMBER: US/11/004,399
CURRENT FILING DATE: 2003-12-03
NUMBER OF SEQ ID NOS: 4043
SOFTWARE: PATENTIN Version 3.1
SEQ ID NO 2080
LENGTH: 11
TYPE: PAT
CORGANISM: SARS-COV Virus
US-11-004-399-2080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; PRIOR FILING DATE: 2003-04-23
; PRIOR APPLICATION NUMBER: 60/465,738
; PRIOR FILING DATE: 2003-04-25
; PRIOR APPLICATION NUMBER: 60/470,935
; PRIOR FILING DATE: 2003-05-14
; NUMBER OF SEQ ID NOS: 2487
; SOFTWARE: PatentIn ver. 3.2
; SEQ ID NO 1387
; LENGTH: 11
; TYPE: PRT
; ORGANISM: Human severe acute respiratory system virus
US-10-807-807-1387
Search completed: May 30, 2006, 15:20:14 Job time : 50.3333 secs
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US-11-004-399-2080
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Best Local Similarity 54.5%;
Matches 6; Conservative
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                                                                                   GOSOSNILPOT 11
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Pred. No. 2.8e+02;
1; Mismatches 4
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Result
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Maximum Match 100%
Listing first 45 s
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score
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     protein search, using
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Match Length
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(without alignments)
50.118 Million cell updates/sec
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Gapop 10.0 ,
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                 7 US-11-219-563-34
7 US-11-254-182-12
7 US-09-731-899-19
6 US-09-731-899-19
6 US-10-522-356-38
7 US-11-252-276-38
7 US-11-252-276-38
7 US-11-258-066-747
7 US-11-19-563-16
7 US-11-177-359C-16
7 US-11-177-359C-16
7 US-11-177-359C-16
7 US-11-177-340C-16
7 US-11-172-986-79
9 US-11-121-286-8
9 US-11-121-286-8
9 US-11-121-282-8
9 US-11-121-282-8
9 US-11-251-7498-53
9 US-11-251-7498-53
9 US-11-267-317-211
9 US-11-140-487A-1314
9 US-11-140-487A-1314
9 US-11-140-487A-1314
9 US-11-140-487A-1314
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Biocceleration Ltd
Sequence 34, Appl Sequence 19, Appl Sequence 19, Appl Sequence 33, Appl Sequence 83, Appl Sequence 82, Appl Sequence 16, Appl Sequence 59, Appl Sequence 59, Appl Sequence 51, Appl Sequence 31, Appl Sequence 31, Appl Sequence 31, Appl Sequence 31, Appl Sequence 3146, Appl Sequence 316, Appl Sequence 316, Appl Sequence 3146, Appl Sequence 5, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Description
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US-11-219-563-34
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| 5                           | 44                 | 43                 | 42                 | 41                | 40               | 9                | 38                | 37               | 36               | <u>ყ</u>         | 34               | 33               | 32                  | 31                | 30                  | 29                 | 28                 | 27                | 26               |
|-----------------------------|--------------------|--------------------|--------------------|-------------------|------------------|------------------|-------------------|------------------|------------------|------------------|------------------|------------------|---------------------|-------------------|---------------------|--------------------|--------------------|-------------------|------------------|
| 20                          | 20                 | 20                 | 20                 | 20                | 20               | 20               | 20                | 21               | 21               | 21               | 21               | 21               | 21                  | 21                | 21                  | 21                 | 21                 | 21                | 22               |
| 20.6                        |                    | 20.6               | ٠                  | 20.6              |                  |                  | 20.6              | 21.6             | 21.6             | 21.6             | 21.6             | 21.6             | 21.6                | 21.6              |                     | 21.6               | ٠                  | 21.6              | 22.7             |
| v                           | φ                  | 9                  | Q                  | v                 | ø                | ø                | v                 | 12               | 11               | 11               | 11               | 11               | 10                  | 10                | Q                   | 9                  | 9                  | 9                 | 17               |
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| US-11-140- <b>4</b> 87A-472 | US-11-140-487A-443 | US-11-140-487A-327 | US-11-140-487A-212 | US-11-140-487A-24 | US-11-140-487A-8 | US-10-538-066-8  | US-10-538-066-7   | US-09-784-950-21 | US-11-177-340C-3 | US-11-177-359C-3 | US-11-177-358C-3 | US-10-540-431-11 | US-11-140-487A-1596 | US-11-055-093-223 | US-11-140-487A-1214 | US-11-140-487A-551 | US-11-140-487A-120 | US-11-055-093-222 | US-11-121-182-52 |
| Sequence 472, F             | •                  | •                  | •                  | Sequence 24, App  | Sequence 8, Appl | Sequence 8, Appl | Sequence 7, Appli | Sequence 21, App | Sequence 3, Appl | Sequence 3, Appl | Sequence 3, Appl | Sequence 11, Ap  | Sequence 1596,      | `                 | 4                   | `                  | -                  | •                 |                  |
| ďď                          | App                | dat                | App                | ρĺ                | 110              | L                | 11                | ğ                | )<br>Li          | 111              | 11               | ğ                | Αp                  | App               | Αp                  | App                | App                | App               | υp1              |

# ALIGNMENTS

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Sequence 34, Application US/11219563

Publication No. US20060088539A1

APPLICANT: Bander, Neil

ITITLE OF INVENTION: MEMBRANE ANTIBODIES TO PROSTATE-SPECIFIC TITLE OF INVENTION: MEMBRANE ANTIBODIES TO PROSTATE-SPECIFIC PILE REFERENCE: 13651.001 (BZL-001)

CURRENT APPLICATION NUMBER: US/11/219,563

CURRENT FILING DATE: 2005-09-02

PRIOR APPLICATION NUMBER: PCT/US04/06586

PRIOR APPLICATION NUMBER: US 10/379,838

PRIOR APPLICATION NUMBER: US 10/379,838

PRIOR APPLICATION NUMBER: 10/449,379

PRIOR APPLICATION NUMBER: 10/449,379

PRIOR FILING DATE: 2003-05-30

NUMBER OF SEQ ID NOS: 144

CONTRADE: FSECTOF OF WINCHOST UNIONS VANCOUS AND MEMBER OF SEQ ID NOS: 144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: PRT ; ORGANISM: Mus musculus US-11-219-563-34
                                                                                                                                                                                                                                                                                                                                      RESULT 2
US-11-254-182-12
APPLICANT: ANDYA, JAMES
APPLICANT: GWEE, SHIANG C.
APPLICANT: LIU, JUN
APPLICANT: SHEN YE
APPLICANT: SHEN YE
TITLE OF INVENTION: ANTIBODY FORMULATIONS
FILE REFERENCE: P2104R1
CURRENT APPLICATION NUMBER: US/11/254,182
CURRENT FILING DATE: 2005-10-19
PRIOR APPLICATION NUMBER: US 60/620,413
                                                                                                                                                                                                                                                  Sequence 12, Application US/11254182 Publication No. US20060088523A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version SEQ ID NO 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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Pred. No. 5
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US-09-731-899-19

Sequence 19, Application US/09731899

; Publication No. US20060088548A1

; GENERAL INFORMATION:

APPLICANT: Chain, Benjamin

; TITLE OF INVENTION: CHIMERIC PEPTIDES AS IMMUNOGENS, ANTIBODIES THERETO, AND

; TITLE OF INVENTION: FOR IMMUNIZATION USING CHIMERIC PEPTIDES OR ANTIBODIES

; FILE REFERENCE: 20555/1203433-US1

; CURRENT FILING DATE: 2000-12-08

; CURRENT FILING DATE: 2000-13-08
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Best Local Similarity 80.0
                                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application US/10522356 Publication No. US20060105323A1 GENERAL INFORMATION:
                                                      SOFTWARE: PatentIn version 3.3
SEQ ID NO 3
LENGTH: 9
TYPE: PRT
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NUMBER OF SEQ ID NOS: 74
SEQ ID NO 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn version 3.3 SEQ ID NO 19
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                                                                                                                                                                                                                      APPLICANT: WHITELAW, CHRISTOPHER BRUCE ALEXANDER APPLICANT: CLARK, ANTHONY JOHN APPLICANT: WOLF, CHARLES ROLAND TITLE OF INVENTION: MULTI-REPORTER GENE MODEL FOR FILE REFERENCE: 102286.155 US1
CURRENT APPLICATION UNMBER: US/10/522,356
CURRENT FILING DATE: 2005.01-26
                                                                                                                            PRIOR APPLICATION NUMBER: PCT/GB03/003192
PRIOR FILING DATE: 2003-07-25
PRIOR APPLICATION NUMBER: GB 0217402.7
PRIOR FILING DATE: 2002-07-26
NUMBER OF SEQ ID NOS: 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/169,687 PRIOR FILING DATE: 1999-12-08
               ORGANISM: Unknown FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
OTHER INFORMATION: Description of Unknown Sequence: Haemaglutinin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Tetanus toxin bacteria
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Pred. No.
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Pred. No.
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red. No. 5.2e+04;
Mismatches 0
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Sequence 83, Application US/11023959A

Dublication No. US20060106203A1

REPLICANT: Winter, Greg
TITLE OF INVENTION: Ligand
FILE REFERENCE: 8039/2142
CURRENT APPLICATION NUMBER: US/11/023,959A
CURRENT FILING DATE: 2004-12-28
PRIOR APPLICATION NUMBER: PCT/GB03/002804
PRIOR APPLICATION NUMBER: PCT/GB03/002804
PRIOR FILING DATE: 2003-06-30
NUMBER OF SEQ ID NOS: 187
SOFTWARE: Patentin version 3.3
SEQ ID NO 83
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Best Local Similarity
Thes 3; Conserve
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Best Local Similarity
Matches 3; Conserv
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                                                                                                                                                                                                                                             US-11-023-959A-83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-11-252-276-38
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LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 1998-10-30
NUMBER OF SEQ ID NOS: 168
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: A System for Cell Base
FILE REFERENCE: 97-022-K2-CO
CURRENT APPLICATION NUMBER: US/11/252,276
CURRENT FILING DATE: 2005-10-17
PRIOR APPLICATION NUMBER: 09/713,572
PRIOR FILING DATE: 2000-11-15
PRIOR APPLICATION NUMBER: 09/430,656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Giuliano, APPLICANT: Bright,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/106,308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/136,078 PRIOR FILING DATE: 1999-05-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 09/031,271
PRIOR FILING DATE: 1998-02-27
PRIOR APPLICATION NUMBER: 08/810,983
PRIOR FILING DATE: 1997-02-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 1999-10-29
PRIOR APPLICATION NUMBER: 09/398,965
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 1999-09-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE: OTHER INFORMATION: Description of Artificial Sequence: HA epitope
                                                                                                                                                                                                                                                                                                                                                              10 FPYTAP 15
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No. US20060094868A1
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Pred. No. 5.2e
1; Mismatches
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Pred. No. 5
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APPLICANT: Bander, Neil
TITLE OF INVENTION: MODIFIED ANTIBODIES TO PROSTATE-SPECIFIC
TITLE OF INVENTION: MEMBRANE ANTIGEN AND USES THEREOF
FILE REFERENCE: 13651.001 (BZL-001)
CURRENT APPLICATION NUMBER: US/11/219,563
CURRENT FILING DATE: 2005-09-02
PRIOR APPLICATION NUMBER: PCT/US04/06586
PRIOR APPLICATION NUMBER: D3/379,838
PRIOR FILING DATE: 2004-03-03
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: D5/03-03
PRIOR APPLICATION NUMBER: D5/03-03-03
PRIOR APPLICATION NUMBER: 10/449,379
PRIOR APPLICATION NUMBER: 10/449,379
PRIOR FILING DATE: 2003-05-30
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; ORGANISM: Artificial sequence
; FEATURE:
; OTHER INFORMATION: HA tag
US-11-023-959A-83
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; OTHER INFORMATION: A3 non-natural consensus peptide
US-10-538-066-747
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Best Local S
Matches 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16, Application US/11219563
Publication No. US20060088539A1
GENERAL INFORMATION:
                                                                                  NUMBER OF SEQ ID NOS: 144
SOFTWARE: FRETSEQ for Windows Version 4.0
SEQ ID NO 16
LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.2 SEQ ID NO 747
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TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7,
TITLE OF INVENTION: Peptides and Compositions
FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR PILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Unknown
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                                      ORGANISM: Artificial Sequence
OTHER INFORMATION: deimmunized light chain J591
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wes 3; Conserv
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Pred. No.
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Pred. No. 1.9e+02;
1; Mismatches 0;
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US-11-177-358C-16

Sequence 16, Application US/11177358C

Publication No. US20060094072A1

GENERAL INFORMATION:
APPLICANT: CAMPBELL, DOUGLAS A.
TITLE OF INVENTION: PEPTIDE SEQUENCE TAGS AND METHOD OF USING SAME FILE REFERENCE: 113703-1006

CURRENT APPLICATION NUMBER: US/11/177,359C

CURRENT FILING DATE: 2005-07-11

NUMBER OF SEQ ID NOS: 16

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 16

LENGTH: 12

TYPE: PRT
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PRIOR FILING DATE: 2002-01-07
PRIOR PELICATION NUMBER: 09/385,219
PRIOR PELICATION NUMBER: 60/318,568
PRIOR APPLICATION NUMBER: 60/118,568
PRIOR FILING DATE: 1999-02-03
PRIOR FILING DATE: 1999-02-03
PRIOR FILING DATE: 1998-08-28
PRIOR APPLICATION NUMBER: 60/098,355
PRIOR FILING DATE: 1998-08-28
PRIOR FILING DATE: 1998-08-28
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Matches
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 82
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Publication No. US20060088846A1
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Best Local Similarity
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TITLE OF INVENTION: METHODS TO IDENTIFY COMPOUNDS USEFUL FOR THE TREATMENT
TITLE OF INVENTION: OF PROLIFERATIVE AND DIFFERENTIATIVE DISORDERS
FILE REFERENCE: 5914-106-999
CURRENT APPLICATION NUMBER: US/11/106,014
CURRENT FILING DATE: 2005-04-13
PRIOR APPLICATION NUMBER: 10/632,150
PRIOR PILING DATE: 2003-07-30
PRIOR FILING DATE: 2003-07-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Pagano, Michele
APPLICANT: Chiaur, Dah Sharim
APPLICANT: Latres, Esther
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/124,449
PRIOR FILING DATE: 1997-03-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
                                            ORGANISM: Artificial Sequence
OTHER INFORMATION: Description of Artificial Sequence: Synthetic peptide
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6; Conserva
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100.0%; Pred. No. 1.9e+02;
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Pred. No.
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Sequence 16, Application US/11177359C

Publication No. US20060094058A1

GENERAL INFORMATION:

APPLICANT: CAMPBELL, DOUGLAS A.

TITLE OF INVENTION: PEPTIDE SEQUENCE TAGS AND METHOD OF USING SAME

FILE REFERENCE: 113703-1004

CURRENT TAPLICATION NUMBER: US/11/177,359C

CURRENT FILING DATE: 2005-07-11

NUMBER OF SEQ ID NOS: 16

SOOTWARE: Patentin Ver. 2.1

SEQ ID NO 16

LENGTH: 12

TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
FEATURE:
COTHER INFORMATION: Description of Artificial Sequence: Synthetic peptide
US-11-177-359C-16
                                                                                                                                                                                                                                                                                                          RESULT 12
US-11-17-340C-16
US-11-177-340C
; Sequence 16, Application US/11177340C
; Publication No. US20060099663A1
; GENERAL INFORMATION:
APPLICANT: CAMPBELL, DOUGLAS A.
; TITLE OF INVENTION: PEPTIDE SEQUENCE TAGS AND METHOD OF USING SAME
FILE REFERENCE: 113703-1005
; CURRENT APPLICATION NUMBER: US/11/177,340C
; CURRENT APPLICATION UNMBER: US/11/177,340C
; CURRENT APPLICATION OF SEQ ID NOS: 16
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 16
; LENCTH: 12
TYPE: PRT
US-11-122-986-799

US-11-122-986-799

; Sequence 799, Application US/11122986

; Publication No. US20060104989A1
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Best Local Similarity
Watches 4; Conserva
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Best Local S
Matches 4
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                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                       Local Similarity tes 4; Conserv
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Pred. No.
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Pred. No. 2.3e+02;
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Pred. No. 2.3e+02;
                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                           Length 12;
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US-11-134-228A-8
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Best Local Similarity
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CURRENT FILING DATE: 2005-05-05
PRIOR APPLICATION NUMBER: 60/423,875
PRIOR FILING DATE: 2002-11-05
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,832
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,915
PRIOR PELICATION NUMBER: 60/423,757
PRIOR FILING DATE: 2002-11-05
PRIOR PILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,758
PRIOR APPLICATION NUMBER: 60/424,367
PRIOR FILING DATE: 2002-11-06
PRIOR FILING DATE: 2002-11-06
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PRIOR FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,370
PRIOR APPLICATION NUMBER: 60/424,362
PRIOR APPLICATION NUMBER: 60/424,362
PRIOR FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,373
PRIOR APPLICATION NUMBER: 60/424,373
PRIOR FILING DATE: 2002-11-06
PRIOR FILING DATE: 2002-11-06
                                                                                                                                                                                                                                   APPLICANT: Lindquist et al.

TITLE OF INVENTION: ELECTRICAL CONDUCTORS AND DEVICES FROM PRION-LIKE PROTEINS
FILE REFERENCE: 30554/40155A

CURRENT APPLICATION NUMBER: US/11/134,228A

CURRENT FILING DATE: 2005-05-20
PRIOR APPLICATION NUMBER: 60/573,277

PRIOR APPLICATION NUMBER: 60/573,277

PRIOR FILING DATE: 2004-05-20

NUMBER OF SEQ ID NOS: 73

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 8

LENGTH: 13

TYPE: PRT
ORGANISM: Artificial Sequence
Query Match
Best Local Similarity 50.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 8, Application US/11134228A Publication No. US20060105320A1 GENERAL INFORMATION:
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SEQ ID NO 799
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APPLICANT: VEDADI, MASOUD
TITLE OF INVENTION: ESSENTIAL NOVEL BACTERIAL POLYPEPTIDES
FILE REFERENCE: IPT-330.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 844
                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Hemagglutinin OTHER INFORMATION: epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 12
TYPE: PRT
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50.0%; Pred. No. 2.5e+02;
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RESULT 15

US-11-257-498-59

US-11-257-498-59

Sequence 59, Application US/11257498

Publication No. US20060088550A1

GENERAL INFORMATION:
APPLICANT: Bachmann, Martin F
APPLICANT: Fulurija, Alma
APPLICANT: Saudan, Phillippe
TITLE OF INVENTION: Gastric Inhibitory Polypeptide (GIP) Antigen Arrays and Used ther
FILE REFERENCE: 1700.0540001

CURRENT APPLICATION NUMBER: US/11/257,498

CURRENT APPLICATION UNMBER: US/11/257,498

CURRENT APPLICATION US/11/257,498

CURRENT APPL
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Search completed: May 30, 2006, 15:20:31 Job time : 4 secs
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              26750, A
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US-09-641-528B-4809
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GENERAL INFORMATION:

APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert

Sequence 4809, Application US/09641528B Patent No. 7026443

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|   | US-08-762-106-17 | US-08-762-106-19 | US-09-700-993-6 | US-10-394-980-55 | PCT-US93-00643-8 | US-09-392-812A-7 | US-08-479-233-8 | US-09-641-528B-4860 | US-09-641-528B-4855 | US-09-573-830-30 | US-09-573-830-4 | US-09-462-645C-29 | US-08-447-515-30 | -08-447-515 | US-08-235-437-30 | -08-235-437 |      |  |
|   | 17, Appl         | 19, Appl         | 6, Appli        | 55, Appl         | 8, Appli         | 7, Appli         | 8, Appli        | 4860, Ap            | 4855, Ap            | 30, Appl         | 4, Appli        | 29, Appl          | 30, Appl         | 4, Appli    | 30, Appl         |             |      |  |

## RESULT 2 US-09-641-528B-4810 밁 Ş US-09-641-528B-4809 TITLE OF INVENTION: USING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS FILE REFERENCE: 2060.0100001 CURRENT PLILOMENTON NUMBER: US/09/641,528B CURRENT FILING DATE: 2000-08-15 PRIOR APPLICATION NUMBER: US 60/172,705 PRIOR APPLICATION NUMBER: US 60/172,705 PRIOR FILING DATE: 1999-12-10 NUMBER OF SEQ ID NOS: 51505 SOFTWARE: FASTSEQ for Windows Version 4.0 SEQ ID NO 4809 Patent No. 7026443 GENERAL INFORMATION: Sequence Best Local Similarity Matches 4; Conserv Query Match APPLICANT: Sette, Alessandro APPLICANT: Sidney, John APPLICANT: Southwood, Scott APPLICANT: Chesnut, Robert APPLICANT: Celis, Esteban APPLICANT: ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Peptide Derived from Human Papillomavirus TYPE: PRT LENGTH: NT: Chesnut, Robert NT: Celis, Esteban NT: Grey, Howard F INVENTION: INDUCING 4810, $\vdash$ 10 5 LIQTDQQATT 14 Sidney, vo.... Southwood, Scott Southwood, Robert Chesnut, Robert Celis, Esteban Grey, Howard MVQVEEQQTT 10 Application US/09641528B Conservative INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS 37.8%; Score 28; DB Pred. No. 70; 4; Mismatches ω --2, Length 10 Indels 0 Gaps 0

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; TITLE OF INVENTION: USING PEPTIDE AND NUCLEI
; FILE REFERENCE: 2060.0100001
; CURRENT APPLICATION NUMBER: US/09/641,528B
; CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
; NUMBER OF SEQ ID NOS: 51505
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 4810
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US-09-641-528B-4810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
TILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 4856
                                                                                                                 Sequence 13626, Application US/09641528B Patent No. 7026443 GENERAL INFORMATION:
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                                             APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
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                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
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ORGANISM: Artificial Sequence
FEATURE:
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Local Similarity 40.0%;
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                   Chesnut, Acc
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Chesnut, Robert
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Pred. No. 78;
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US-09-641-528B-47147
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                                                                                                      Sequence 47147, Application US/09641528B Patent No. 7026443 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 26703
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Best Local
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Best Local Similarity 40.0%;
                                                                                                                                                                                                                                                                                                                                         Matches
APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esteban
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esteban
APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REPERENCE: 2060.0100001
FULE REPERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR PILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
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APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
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OTHER INFORMATION: Peptide Derived
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Peptide Derived from Human Papillomavirus
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4; Conservative
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Pred. No.
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CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 50653
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE.
                                                                                                   RESULT 8
US-09-641-528B-4922
; Sequence 4922, Application US/09641528B
; Patent No. 7026443
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; Sequence 50653, Application US/09641528B
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
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CURRENT APPLICATION NUMBER: US/09/641,528B

CURRENT FILING DATE: 2000-08-15

PRIOR APPLICATION NUMBER: US 60/172,705

PRIOR FILING DATE: 1999-12-10

NUMBER OF SEQ ID NOS: 51505

SOFTWARE: PASTSEQ for Windows Version 4.0

SEQ ID NO 47147
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Best Local
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APPLICANT:
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APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Sette, Alessandro APPLICANT: Sidney, John
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Peptide Derived from Human Papillomavirus -09-641-528B-47147
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ORGANISM: Artificial Sequence
FEATURE:
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Local Similarity 40.0%;
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; Pred. No. 1.1e
4; Mismatches
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Pred. No. 1.1e+02;
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RESULT 10 US-09-641-528B-4923

Sequence 4923, Application US/09641528B Patent No. 7026443
GENERAL INFORMATION:
APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott

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                                                                                                            ; OTHER INFORMATION: Peptide Derived from Human Papillomavirus US-09-641-528B-50984
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Best Local S
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Best Local
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APPLICANT:
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CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Sette, Alessandro APPLICANT: Sidney, John APPLICANT: Southwood, Scott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
                                                                                                                                        ORGANISM: Artificial Sequence FEATURE:
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                                                 Local Similarity les 4; Conserv
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Celis, Esteban
Grey, Howard
                                                    Conservative
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                                                                 Score 26;
Pred. No.
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Pred. No. 5e+05;
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5e+05;
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RESULT 12
US-09-236-415-6
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Sequence 6, Application US/09236415
PAtent NO. 6309863
GENERAL INFORMATION:
APPLICANT: Anderson, Carl W.
APPLICANT: Appella, Ettore
                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 26799
LENGTH: 10
TYPE: PRT
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APPLICANT: Grey, Howard
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR TILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4923
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 26799, Application US/09641528B Patent No. 7026443
                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
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Local Similarity 44.4%;
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                                                                                                                                                                                                                                                                                                   35.1%;
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Pred. No. 1.6e+02;
3; Mismatches 2
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Pred. No. 1.6e+02;
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                                                                                                                                                                                                                                                                                                                         DB 3; Length 10;
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US-08-221-583-42
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US-08-221-583-42
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                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 42, Application US/08221583
Patent No. 5486595
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 6
    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                    TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAMME: DeLuca, Mark
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: CCOR
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Sakaguchi, Kazuyasu
TITLE OF INVENTION: METHODS FOR GENERATING PHOSPHORYLATION SITE-SPECIFIC
TITLE OF INVENTION: IMMUNOLOGICAL REAGENTS
FILE REFERENCE: U.S. Application 09/236,415
CURRENT APPLICATION NUMBER: US/09/236,415
CURRENT FILLING DATE: 1999-01-25
CURRENT FILLING DATE: 1999-01-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Residue Xaa is a phosphoserine mimetic
                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25:mdctcMod.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/221,583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Tumor Necrosis Factor Inhibitors NUMBER OF SEQUENCES: 62
CORRESPONDENCE ADDRESS:
ADDRESSE: Woodcock Washburn Kurtz Mackiewicz & No. STREET: One Liberty Place 46th Floor CITY: Philadelphia
STATE: Pennsylvania
                                                                                                MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH: 14
Local Similarity tes 5; Conserv
                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
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    Conservative
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                      33.8%;
55.6%;
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54.5%;
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Pred. No. 3.6e+02;
                      Score 25; DB 1;
Pred. No. 3.9e+02;
      Mismatches
                                        Length 15;
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PCT-US95-04018-42 RESULT 14

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PCT-US95-04018-42
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Best Local S
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                                                                               Patent No. 5932415
GENERAL INFORMATION:
                                                                                                 Sequence 38, Application US/08456670B Patent No. 5932415
                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/221,580
FILING DATE: 01-APR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/221,583
FILING DATE: 01-APR-1994
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Deluca, Mark
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: CCC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 0 FILING DATE: 01-APR-1994 ATTORNEY/AGENT INFORMATION:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Heavner, George A. APPLICANT: Kruszynski, Marian
                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                ELECUTATION (215)
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                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
TOPOLOGY: linear
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Similarity 55.6%;
5; Conservative
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SCHUBERT, PETER
NEUWANN, SIEGFRIED
PAWELZIK, MARTINA
LINXWEILER, WINFRIED
BURGER, CHRISTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mervic, Miljenko
Weber, Robert W.
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Mervic, Miljenko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Woodcock Washburn Kurtz Mackiewicz &
                                                                                                                                                                                                                                                                                                                                      peptide
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Pred. No. 3.
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3.9e+02;
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Search completed: May 30, 2006, 11:11:06 Job time : 20.3333 secs .

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US-08-456-670B-38
                                                                  Query Match
Best Local S
Matches 4
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CLASSIFICATION: 435

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/412,227

FILING DATE: 27-MAR-1995

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/075,248

FILING DATE: 11-JUN-1993

PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE 4239567.4

FILING DATE: 25-NOV-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/456,670B
                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE 421
APPLICATION NUMBER: DE 421
FILING DATE: 11-UUM-1992
ATTORNEY/AGENT INFORMATION:
NAME: HAMLET-KING, DIANA
REGISTRATION NUMBER: 33,34
REFERENCE/DOCKET NUMBER: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: MILLEN, WHITE, ZELANO & BRANIGAN, P.C.
STREET: 2200 CLARENDON BLVD., SUITE 1400
                                                                                                                                                                            ORIGINAL SOURCE:
                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 703-243-6333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                       TELEPHONE: YUS-1-1010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: ARLINGTON
STATE: VIRGINIA
                                                                                                                                                                                                                                                                                                  TELEX:
                                                                                Local Similarity
                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                            TYPE: amino acid
                                                                                                                                                            ORGANISM:
EQQTTTK 7
                                 DQQATTR 15
                                                                                                                                                                                                                                                                                                     64191
                                                                                                                                                                                                                                                  11 amino acids
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GOEBEL, WERNER
KOHLER, STEFAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HOFMANN,
                                                                  32.4%;
llarity 57.1%;
Conservative
                                                                                                                                                          Listeria innocua
                                                                                                                                                                                                               linear
                                                                                                                                                                                         peptide
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                                                                                    Score 24; DB 1;
Pred. No. 4.2e+02;
                                                                     Mismatches
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Result
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Perfect score:
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Maximum Match
Listing first
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Maximum DB seq length: 15
   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                           Score
   %
Query
Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       May 30, 2006, 09:58:55; Search time 11.6667 Seconds (without alignments) 123.707 Million cell updates/sec
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deoxynucleotidyltr
Ig heavy chain V r
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Ig kappa chain V-I
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| g &                            | 3 m 0                                | RESULT 2 B61458 Ig kappa chain V-I re C;Species: Homo sapit C;Date: 15-Oct-1994 † C;Accession: B61458 R;Brouet, J.C.; Dellit J. Exp. Med. 170, 155 A;Title: Expression A;Reference number: A;Reference number: A;Reference number: B61458 A;Status: preliminary A;Molecule type: prot A;Refores: 1-13 <-BRC A;Cross-references: UC;Keywords: heterotet  | 유양                                    | 3 m O                                 | A;Reference number: S<br>A;Accession: S32677<br>A;Status: preliminary<br>A;Molecule type: DNA<br>A;Residues: 1-15 <mon<br>A;Cross - references: U<br/>C;Superfamily: nitros<br/>C;Keywords: sulfurtra</mon<br>   | R; A   | C; G; C; C; S; C; C; S; C;              |            |  |
|--------------------------------|--------------------------------------|---|---------------------------------------|---------------------------------------|--|--|---|------------|--|
|                                | Query Ma<br>Best Loo<br>Matches      | RESULT 2 B61458 Ig Kappa chain V-I C;Species: Homo sap C;Date: 15-Oct.1994 C;Accession: B61458 I, Recession: B61458 I, Tille: Expression A;Reference number: A;Accession: B61458 A;Status: prelimina A;Molecule type: pr A;Reference type: pr A;Residues: 1-13 <b 1-13="" <b="" a;residues:="" heterence<="" td=""><td></td><td>Query Ma<br/>Best Loo<br/>Matches</td><td>efere<br/>ccess<br/>tatus<br/>olecu<br/>esidu<br/>ross-<br/>uperf</td><td>;ACCESSION: 8326//<br/>;Monnerjahn, U.; B<br/>ubmitted to the EM<br/>;Description: Clon</td><td>RESULT 1 S32677 nitrogenase co N;Contains: L- C;Species: Ana C;Date: 06-Jan</td><td></td><td>0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0</td></b>   |                                       | Query Ma<br>Best Loo<br>Matches       | efere<br>ccess<br>tatus<br>olecu<br>esidu<br>ross-<br>uperf  | ;ACCESSION: 8326//<br>;Monnerjahn, U.; B<br>ubmitted to the EM<br>;Description: Clon | RESULT 1 S32677 nitrogenase co N;Contains: L- C;Species: Ana C;Date: 06-Jan |            | 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0  |
|                                | Match<br>Local<br>les                | n chass: H<br>15-O<br>15-O<br>15-O<br>15-O<br>15-O<br>15-O<br>15-O<br>15-O  |                                       | Match<br>Local<br>Les                 | ion: i pr ile t les: refe amil   | jahn<br>d to   | nase cofacto<br>ins: L-cyste<br>es: Anabaena<br>06-Jan-1927                 |            |  |
| 6<br>10<br>0                   | b<br>Simi<br>5;                      | chain V-I re Homo sapie 5-Oct-1994 # 5-Oct-1994 # On: B61458 O.C.; Della Oct. Della Con Bxyression o Cxyression o Cn B61458 Preliminary Expression o Con B61458 Oct. B6105 Oct. B705 Oct. | 5 LI                                  | h<br>Simi<br>5;                       | nce number: S3 ion: S32677 : preliminary : preliminary DNA es: 1-15 <mon: amily:="" ds:="" nitroge="" references:="" sulfurtrar<="" td="" un=""><td>the</td><td>cofact<br/>L-cyst<br/>Anabaer<br/>Jan-199</td><td></td><td>1155<br/>1166<br/>1166<br/>1166<br/>1166<br/>1166<br/>1166<br/>1166</td></mon:> | the  | cofact<br>L-cyst<br>Anabaer<br>Jan-199                                      |            | 1155<br>1166<br>1166<br>1166<br>1166<br>1166<br>1166<br>1166   |
| QTDQQATT<br>      <br>QMTQSPTT | ilarity<br>Conser                    | 7-I region sapiens 994 #sequ. 458   1551-11   1551-12   1501 of a sion of a | ALDOL<br> <br>DOLLO                   | ilarit;<br>Conse:                     | Aber: S32<br>32677<br>3: DNA<br>3: DNA<br>15 <mon><br/>10ces: UNI<br/>1nitrogen<br/>1nitrogen</mon>  | U.; Boehme,<br>the EMBL Dat<br>Cloning an  | <b>□. □</b>   |            | 2200 - 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6   |
| <b>_</b>                       | rva                                  | in V-I region (BL) como sapiens (man) ct-1994 #sequence, B61458 C.; Dellagi, K.; 170, 1551-1558; ression of a publ. number: A61458; M B61458 ellminary ype: protein 1-13 <bro> rences: UNIPARC:U</bro>  | LIQTDQQATTR :           : VIYLDNNATTK | .ty<br>serva                          | in p   | U.; Boehme, H<br>the EMBL Data<br>1: Cloning and                                     | synthesi<br>ne sulfur<br>variabili<br>#sequence                             |            |  |
| 4 0                            | 28.<br>55.<br>tive                   | (BI<br>, who  | 15                                    | 33.8<br>Y 45.5<br>rvative             | S32675  CY  NN> UNIPROT:Q445  gen fixation cansferase  | H.<br>a Lik<br>d exp   | synthesis protein ne sulfurtransferas variabilis #sequence_revision         |            | 10<br>10<br>11<br>11<br>11<br>11<br>11<br>11<br>11<br>11<br>11<br>11<br>11<br>1  |
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|                                | Sco<br>Pre<br>0;                     | A) - human (f<br>revision 15-<br>Gendron, M.C.<br>1989<br>ic idiotype b<br>IUID:90039128;   |                                       | Score<br>Pred.<br>2; Mis              | 507; UNIPA   | H.<br>Library, De<br>expression  | otein   | Þ          | D54823<br>S70720<br>A33099<br>E828425<br>E906681<br>P706681<br>P4163<br>P41636<br>A45096<br>A35527<br>A35527<br>A35527<br>P507682<br>P707682   |
|                                | Score 21;<br>Pred. No.<br>0; Mismato | m 15<br>m 15<br>M.C<br>ype J<br>9128  |                                       | Mism                                  | NIPA<br>ein  | in   | 96 e p.   | LIGN       | 8880<br>909<br>1944<br>1990<br>1990  |
|                                | <del>5</del> -                       | Y ~ O H   |                                       | re 25; DB<br>d. No.·2.1<br>Mismatches | RC:U   |  | fs - )<br>(EC 2   | ALIGNMENTS |  |
|                                | DB 2;<br>e+03;<br>es                 | (fragment) 5-OGt-1994 C.; Cheval by human   by human   by PMID:24 8; PMID:24  |                                       | ; DB 2; I<br>.·2.1e+02;<br>cches 4;   | 00014  | : 1992<br>coli o   | Anab<br>2.8.1   | vs         |  |
|                                | 4.                                   | agment)<br>Gt-1994 #text<br>Chevalier, A<br>human monocl<br>PMID:2478651  |                                       | ; Le<br>52;<br>4;                     | UNIPARC:UPI00001301CC  | tt.  | # - )<br>ae n   |            |  |
|                                | Length 1; Indel                      | Ë ≯ ë   |                                       | Length; Ind                           | )100;  | he An  | ×   |            |  |
|                                | 13;<br>els                           | 0 5   |                                       | gth 15;<br>Indels                     |  | Anabaena   | variabilis<br>t_change 0  |            | olfi<br>cri<br>1633<br>20K<br>pro<br>pho<br>32K<br>juv<br>juv<br>juv<br>juv<br>pho<br>call<br>alc<br>hyp   |
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|                                | G<br>G                               | 16-Aug-199<br>tt, C.; Mi  |                                       |                                       | 8686   |  | (frag   |            | ttory receptiory receptory receptor factor receptor factor excentigen rottein - rottein 30 rottein 30 rottein 30 rottein - rottein 30 rottein 30 rottein 30 rottein 30 rottein 30 rottein 30 rottein 90 rottein 9 |
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|                                |                                      | myelin-asso   |                                       |                                       | NID: 9296503   |  |   |            |  |
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A; Genome: nu
C; Keywords:
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Eur. J. Biochem. 221, 1111-1116, 1994
A;Title: Identification of tissue-specific
A;Reference number: S43624; MUID:94237150;
                                                                                                                                                                                                                                                                                                                                       cytochrome-c oxidase (EC 1.9.3.1) chain Va, hepatic - rainbow trout (fragment) C;Species: Oncorhynchus mykiss (rainbow trout) C;Date: 20-Oct-1994 #sequence_revision 01-Nov-1996 #text_change 09-Jul-2004
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A; Residues: 1-15 < WAS>
                                                                                                                                                            A; Molecule type: protein A; Residues: 1-10 < FRE>
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C;Date: 15-Oct-1994 #sequence revision 15-Oct-1994 #text_change 16-Aug-1996
C;Accession: A61458; PLOIS6
R;Brouet, J.C.; Dellagi, K.; Gendron, M.C.; Chevalier, A.; Schmitt, C.; Mihaesco, E.
J. Exp. Med. 170, 1551-1558, 1989
A;Title: Expression of a public idiotype by human monoclonal IgM directed to myelin-assc A;Reference number: A61458; MUID:90039128; PMID:2478651
A;Accession: A61458
                                                                                                                  A; Note:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, J. Exp. Med. 176, 1577-1581, 1992
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C;Comment: This protein is one of monoclonal
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  Query Match
                                                                                          Genetics:
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;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
                                                                       Genome: nuclear
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Similarity 55.6%;
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is designated as
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Salmo gairdneri
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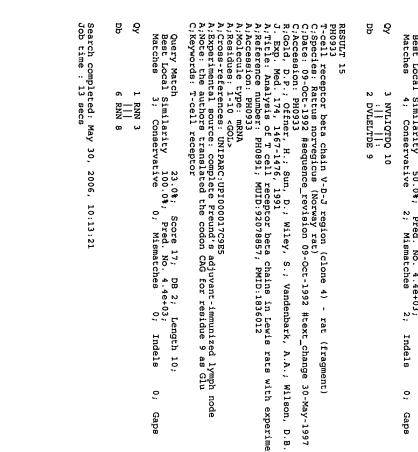
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urinary tract stone matrix protein 10, 42K - human (fragment) C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 12-Apr-1995 #sequence_revision 12-Apr-1995 #text_change C;Accession: H56046 W.B.; Gawinowicz, M.A.; Kendrick, N. g;Binette, J.P.; Binette, M.B.; Gawinowicz, M.A.; Kendrick, N. submitted to the Protein Sequence Database, Fabruary 1995 A;Description: Isolation, characterization and sequence of stone
                                                                                                                                                                synaptosomal-associated protein SNAP-25 peptide 8 - rabbit (fragment)
N,Alternate names: superprotein peptide
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 31-Mar_1993 #sequence_revision 18-Nov-1994 #text_change 15-Jun-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R.Lark, M.W.; Gordy, J.T.; Weidner, J.R.; Ayala, J.; Kimura, J.H.; Williams, J. Biol. Chem. 270, 2550-2556, 1995
A;Title: Cell-mediated catabolism of aggrecan. Evidence that cleavage at the A;Reference number: A55695, MUID:95155314; PMID:7852317
A;Accession: A55695
                                                              R;Loewy, A.; Liu, W.S.; Baitinger, C.; Willard, M.B.
J. Neurosci. 11, 3412-3421, 1991
A;Title: The major 35S-methionine-labeled rapidly transported
A;Reference number: A44823; MUID:92044785; PMID:1941090
  A; Molecule type:
                       A; Status: preliminary
                                           A; Accession: C44823
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A;Cross-references: UNIPROT:Q7M4P7; UNIPARC:UPI000017C406
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A;Residues: 1-10 <LAR>
A;Cross-references: UNIPARC:UPI000017C9A9
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C;Species: Ra
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A;Residues: 1-13 <BIN>
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Rattus norvegicus (Norway rat)
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Pred. No. 1.8e+03;
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A;Residues:

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RESULT 10
S43626
          epithelial dog allergen - Canis familiaris (fragment)
C;Specites: Canis familiaris
C;Date: 27-Oct-2003 #sequence_revision 27-Oct-2003 #text_change 27-Oct-2003
C;Accession: A59491
R;Saarelainen, S.; Taivainen, A.; RytkM_vnen_Nissinen, M.; Auriola, S.; Imm
                                                                                                                                                                                                                                                                                                                                                                                   Eur. J. Biochem. 221, 1111-1116, 1994
A;Title: Identification of tissue-specific
A;Reference number: $43624; MUID:94237150;
A;Accession: $43626
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytochrome-c oxidase (EC 1.9.3.1) chain Vb-H - trout (fragment) C;Species: Salmo sp. (trout) C;Date: 19-Mar-1997 #sequence_revision 01-Aug-1997 #text_change C;Accession: S43626 R;Freund, R;Kadenbach, B.
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S66636
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A;Experimental source: visual tissue
A;Note: sequence extracted from NCBI back
C;Keywords: membrane trafficking
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C;Keywords: electron transfer; membrane-associated complex; oxidoreductase; respiratory
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A;Title: Crystallisation and preliminary X-ray analysis of the receptor-binding domain A;Reference number: S66634; MUID:96032553; PMID:7556651
A;Accession: S66636
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                                                                                                      A59491
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A;Residues: 1-11 <FRE>
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C;Species: Bos primigenius indicus (zebu cattle)
C;Date: 19-Mar-1997 #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: UNIPROT:Q7M2N7; UNIPARC:UPI000017C480
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Pred.
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Pred. No.
RytkM-vnen-Nissinen, M.; Auriola, Database, October 2003
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3.2e+03;
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2.7e+03;
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RESULT 13
S72432
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Blochem. J. 319, 499-506, 1996
A;Title: purification and characterization
A;Reference number: $72431; MUID:97069704;
A;Accession: $72432
A;Molecule type: protein
A;Residues: 1-15 <CHA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: Homo sapiens (man)
C;Date: 23-Nov-1991 #sequence_revision 23-Nov-1991 #text_
C;Dates: 23-Nov-1991 #sequence_revision 23-Nov-1991 #text_
C;Accession: PT0205; A31175
R;Bautista, C.M.; Baylink, D.J.; Mohan, S.
Biochem. Biophys. Res. Commun. 176, 756-763, 1991
Biochem. Biophys. Res. Commun. 176, 756-763, 1991
A;Title: Isolation of a novel insulin-like growth factor
A;Reference number: PT0205; MUID:91222244; PMID:1709017
A;Accession: PT0205
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PT0205
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A;Accession: A59491
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                                                                                                                                 A;Description: NADP-dependent lipoamide reductase A;Pathway: epoxypropane degradation A;Note: sensitive to N-ethyl-maleimide and p-chloromercuribenzoate C;Keywords: FAD
                                                                                                                                                                                                                                       A;Cross-references: UNIPARC:UPI000017CA91A;Note: 3-Met and 13-Leu were also found
                                                                                                                                                                                                                                                                                                                                                                                          A; Variety: strain Py2
C; Date: 14-Apr-1998 #sequence_revision
C; Accession: $72432
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C; Species: Xanthobacter sp.
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                                                                Query Match
Best Local S
Matches 4
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33.3%;
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Pred. No.
                                                                 Score 18; DB 2;
Pred. No. 4.5e+03;
2; Mismatches
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4.5e+03;
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RESULT 15
PH0933
C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997
C;Accession: PH093
R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, S.; Vandenbark, A.A.; Wilson, D.B.
J. Exp. Med. 174, 1467-1476, 1991
A;Title: Analysis of T cell receptor beta chains in Lewis rats with experimental allergi A;Reference number: PH0991; MUID:92078857; PMID:1836012
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A61354
A61354
Carnitine medium/long chain acyltransferase (EC 2.3.1.-) - rat (fragment)
Carnitine medium/long chain acyltransferase (EC 2.3.1.-) - rat (fragment)
N,Alternate names: endoplasmic reticulum protein ERp61; glucose regulated protein GRP58;
C;Species: Rattus norvegicus (Norway rat)
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 07-May-1999
C;Accession: A61354
R;Murthy, M.S.R.; Pande, S.V.
Mol. Cell. Blochem. 122, 133-138, 1993
A;Title: Carnitine medium/long chain acyltransferase of microsomes seems to be the previ A;Accession: A61354
A;Accession: A61354
A;Accession: A61354
A;Accession: A61354
A;Status: preliminary
A;Accession: A61354
Search completed: May 30, 2006, 10:13:21 Job time : 13 secs
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                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPARC:UPI000017C9E5
A;Experimental source: complete Freund's adjuvant-immunized lymph node
A;Note: the authors translated the codon CAG for residue 9 as Glu
C;Keywords: T-cell receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: mRNA
A; Residues: 1-10 <GOL>
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Best Local S
Matches 3
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Q5TRH2_GOVIN
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Q9UJ48 HUWAN
Q6MZZ1 HUWAN
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Q5GJB2 COXBU
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DT 05-JUL-2005, sequence version 1.
DT 07-FEB-2006, entry version 4.
DE Hypothetical protein (Fragment).

GN ORFNames=PC106890.00.0;
OS Plasmodium chabaudi.
CN CHARYOta; Alveolata; Apicomplex
NCBI_TaxID-5825;
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RA Guail M.A., Ormond D., Doggett J.
RA Gafacos F.C., Janse C.J., Barrel
RA Sinden R.S.;
RT "A comprehensive survey of the P
RT Transcriptomic, and proteomic an
Science 307:82-86 (2005).
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                                                                                                                              transcriptomic, and proteomic analyses.";
Science 307:82-86(2005).
-!- CAUTION: The sequence shown here is derived.
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MEDLINE=96073661; PubMed=8540346;

Faro C., Verissimo P., Lin Y., Tang J., Pir
Farotosin A and B. aspartic proteases from
Adv. Exp. Med. Biol. 362:373-377(1995).
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RESULT 5
Q9MZS4 C
AC Q9MZS4;
AC Q9MZS4;
DT 01-CCT-2
DT 01-CCT-2
DT 07-FEB-2
DE Growth h
OS Canis fa
OC Enkaryot
OC Mammalia
OC Canis
OC CANIS
OX NCBI_Tax
RN NUCLEOTI
RX MEDLINE=
RA Lantinga
RT growth h
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ID P70740_1
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Best Local S
Matches 4
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Best Local S
Matches 4
                                                                                             Canis familiaris (Dog).
Canis familiaris (Dog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Carnivora; Fissipedia; Canida
                                                                                                                                                                                                                                                                                                                                                                                             EMBL; XT
NUCLEOTIDE SEQUENCE.

MEDILINE=2231938; PubMed=12431806; DOI=10.1016/S0303-7207(02)00257-5;
Lantinga-van Leeuwen I.S., Timmermans-Sprang E.A., Mol J.A.;

"Cloning and characterization of the 5'-flanking region of the canine growth hormone gene.";
                                                                                                                                                       01-OCT-2000, integrated into UniProtKB/TrEMBL 01-OCT-2000, sequence version 1. 07-FEB-2006, entry version 8.
                                                                                                                                                                                               Q9MZS4_CANFA
Q9MZS4;
                                                                                                                                                                                                                                                                                                                                                                                                                            Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
                                                                                                                                           Growth hormone (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       J. Bacteriol. 176:4385-4393(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=94292469; PubMed=8021224; Lenz O., Schwartz E., Dernedde J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ralstonia eutropha (Alcaligenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Name=hoxA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HoxA protein
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01-FEB-1997,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "The Alcaligenes eutrophus
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SDKQAT
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Betaproteobacteria; Burkholderiales;
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Pred. No. 4.1e+03;
1; Mismatches 1
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Pred. No. 2.
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X gene participates in h
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Maruyama
               PROTEIN SEQUENCE.
MEDLINE=93250579; PubMed=8387370;
                                                               NCBI_TaxID=9913;
                                                                                    Pecora; Bovidae; Bovinae;
E., Iwamatsu
                                                                                     Вов
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RESULT 7
GOTTRI GOTTRI
ID GOTTRI
AC QOTTR
DT 01.-M
DT 01.
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SEQUENCE
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OTRQ7;
01-MAY-2000, integrated into Unil
01-MAY-2000, sequence version 1.
07-FEB-2006, entry version 9.
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01-MAY-2000, sequence version 1.
07-FEB-2006, entry version 11.
Alkaline phosphodiesterase I (EC 3.1.4.1) (Fragment).
Bos taurus (Bovine).
                                                                                                                                                                                                                                                                                                          Q9TRH1_BOVIN
Q9TRH1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia;
Pecora; Bovidae; Bovinae; Bos.
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                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "CAP-50, a newly identified annexin, fibroblast 3Y1 cells.";
J. Biol. Chem. 267:13498-13504(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mizutani A., Usuda N., T
Kobayashi R., Hidaka H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=92317074;
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NON_TER 12 12
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66.7%;
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Pred. No. 6.4e
1; Mismatches
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Pred. No. 5.4e+03;
2; Mismatches (
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Attribution-NoDerivs License
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RESULT 9
Q75MV9 HUMAN
ID Q75MV9 HUMAN
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   RRR OCCOON DETEN
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Q6JDM9_CANFA
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Best Local
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QGJDM9;
05-JUL-2004,
05-JUL-2004,
07-FEB-2006,
                                                                                                          05-JUL-2004,
05-JUL-2004,
21-FEB-2006,
Hypothetical
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NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                           PubMed=15233990; DOI=10.1016/j.ygeno.2004.04.001; Housley D.J.E., Ritzert E., Venta P.J.; "Comparative radiation hybrid map of canine chromosome incorporating SNP and indel polymorphisms."; Genomics 84:248-264(2004).
                                                                                                                                                                                                                                                                                                                                                                                     Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Canis familiaris (Dog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Laurasiatheria; Carnivora; Fissipedia; Canidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUL-2004, sequence version 1.
07-FEB-2006, entry version 8.
Mitochondrial H+ transporting ATP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GO; GO:0004528; F:phosphodiesterase I activity; IEA. SEQUENCE 15 AA; 1678 MW; BE31498EEA8B7071 CRC64;
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                                                                      Homo sapiens (Human)
Eukaryota; Metazoa;
                                                                                                 Name=SSBP1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Name=ATP5A1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biochem. Mol. Biol. Int. 29:579-586(1993).
NUCLEOTIDE SEQUENCE.
MEDLINE=22737999; PubMed=12853948; DOI=10.1038/nature01782;
                                  NCBI_TaxID=9606;
                                                  Homo
                                                             Mammalia;
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                                                            Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
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50.0%;
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Pred. No. 6.
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Pred. No..6.
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RA Wagner-McPherson C., Layman D., Maas J., Jaeger S., Walker R.,
RA Wylie K., Sekhon M., Becker M.C., O'Laughlin M.D., Schaller M.E.,
RA Fewell G.A., Delehaunty K.D., Miner T.L., Nash W.E., Cordes M., Du H.,
RA Sun H., Edwards J., Bradshaw-Cordum H., Ali J., Andrews S., Isak A.,
RA Vanbrunt A., Nguyen C., Du F., Lamar B., Courtney L., Kalicki J.,
RA Czersky P., Bielicki L., Scott K., Holmes A., Harkins R., Harris A.,
RA Strong C.M., Hou S., Tomlinson C., Dauphin-Kohlberg S.,
RA Strong C.M., Hou S., Tomlinson C., Dauphin-Kohlberg S.,
RA Kozlowicz-Reilly A., Leonard S., Rohlfing T., Rock S.M.,
RA Latreille P., Miller N., Johnson D., Murray J., Woessner J.P.,
RA Hickenbotham M.T., Eldred J., Williams D., Bedell J.A., Mardis E.R.,
RA Bieri T.A., Nelson J.O., Berkowicz N., Wohldmann P.E., Cook L.L.,
RA Hickenbotham M.T., Eldred J., Williams D., Bedell J.A., Mardis E.R.,
RA Gillett W., Zhou Y., James R., Phelps K., Iadanoto S., Bubb K.,
RA Gillett W., Zhou Y., James R., Phelps K., Iadanoto S., Bubb K.,
RA Simms E., Levy R., Clendenning J., Kaul R., Kent W.J., Furey T.S.,
RA Balley J.A., Portnoy M.E., Torrents D., Chinwalla A.T., Gish M.R.,
RA Balley J.A., Portnoy M.E., Torrents D., Chinwalla A.T., Gish M.R.,
RA Eddy S.R., McPherson J.D., Olson M.V., Eichler E.E., Green E.D.,
RT "The DNA scenerace of human chromosome 7":
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OSTRITR
PRELIMINARY; PRT; 14 AA.
P83330;
O1-JUN-2002, integrated into UniProtKB/TrEMBL.
O1-JUN-2002, serguence version 1.
O7-FEB-2006, entry version 8.
O7-FEB-2006, entry version 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      M protein, serotype 32 (Fragment). Streptococcus thermophilus.
                                                                                                                                                                Streptococcus thermophilus grown Lait 0:0-0(2002).
                                                                                                                                                                                                                                                                    Guimont C.
                                                                                                                                                                                                                                                                                           STRAIN-ITGST82;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacteria; Firmicutes;
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Submitted
                                                                                                                                                                                          Juimont C., Chopard M.A., Gaillard J.L., Chamba J.F.; "Comparative study of the protein composition of three strains of Streptococcus thermophilus grown either in M17 medium or in milk.";
                                                                                                                                                                                                                                                                                                                             PROTEIN SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hypothetical protein.
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   of protein M. Protein M is closely associated with the bacterium and can render the organism resistant phagocytosis (By similarity).
SUBCELLULAR LOCATION: Type I membrane protein. Cell
                                                                                                FUNCTION: This protein is one of the different antigenic serotypes of protein M. Protein M is closely associated with virulence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 3; Conserv
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37.5%;
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   Type I membrane protein. Cell wall
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J.2e+03;
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RESULT 11
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GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0009405; P:pathogenesis; IEA.
GO; GO:0006909; P:phaggocytosis; IEA.
Antigen; Cell wall; Phagocytosis; Transmembrane; Virulence.
NON_TER 1 1
NON_TER 1 1 1
SEQÜENCE 14 AA; 1739 MW; 66339B9F3A3FE24B CRC64;
                                                                                                                                                                                                                                                                    Fuschiotti P., Harindranath N., Mage R.G., McCormack W.7 Dhanazajan P., Roux K.H.; "Recombination activating genes-1 and -2 of the rabbit: characterization of germline and expressed genes."; Mol. Immunol. 30:1021-1032(1993).
                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE.
STRAIN=RIM/rgm; TISSUE=Thymus;
MEDLINE=93354283; PubMed=8350872; DOI=10.1016/0161-5890(93)90127-W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUL-2004, sequence ver 07-FEB-2006, entry versic Recombination activating
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                                                                                                       Pfam; PF03089; RAG2; 1.
NON TER 15 15
SEQUENCE 15 AA; 1717 MW;
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                                                                                                                                                                                                                              Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                         Oryctolagus.
NCBI_TaxID=9986;
                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oryctolagus cuniculus (Rabbit).
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                                                                                                                                                   InterPro;
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SIMILARITY: TO OTHER STREPTOCOCCAL AND STAPHYLOCOCCAL PROTEINS
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                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                       Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Glires; Lagomorpha; Leporidae;
                                                                 28.4%;
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                                                                 Score 21; DB 2;
Pred. No. 1.1e+04;
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Pred. No. 1e+04;
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                                                                             Length 15;
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RESULT 12 Q9TR62\_RABIT

Q9TR62; 01-MAY-2000, Q9TR62\_RABIT

integrated into

UniProtKB/TrEMBL.

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01-OCT-1994, i
01-OCT-1994, s
07-FEB-2006, (
                                                                                                                                                                                                                                                                                                                                                    "Identification of tissue-specific isoforms for subunits Vb and VIIa of cytochrome c oxidase isolated from rainbow trout.";
Eur. J. Biochem. 221:111-1116(1994).
-!- FUNCTION: This is the heme A-containing chain of cytochrome c oxidase, the terminal oxidase in mitochondrial electron transport.
-!- CATALYTIC ACTIVITY: 4 ferrocytochrome c + O(2) = 4 ferricytochrome
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07-FEB-2006, entry version 9.
Apolipoprotein A.-IV (Fragment).
Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Actinopterygii; Neopterygii; Teleostei; Euteleostei Protacanthopterygii; Salmoniformes; Salmonidae; Onco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Genetic but not diet-induced hypercholesterolemia causes apolipoprotein A-IV level in rabbit sera."; Atherosclerosis 113:171-178(1995).
NON TER
SEQUENCE
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                                                                                                                               Direct
                                                                                                                                                       PIR; S43625;
                                                                                                                                                                                                          Distributed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=8022;
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ONCMY
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Mezdour H., Yamamura T., Nomura S., Yamamoto A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROTEIN SEQUENCE.
                                                                                                  Metal-binding;
                                                                                                                                                                                                                                                                          SUBCELLULAR LOCATION: mitochondrion; mitochondrial SIMILARITY: Belongs to the cytochrome c oxidase Va
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                                                                                               protein sequencing; Heme; Inner membrane; Iron; binding; Mitochondrion; Oxidoreductase.
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Pred. No. 1.1e+04;
  C535C5B1AB02C33D CRC64;
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RESULT 14
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DT 07-FEB-2
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ID Q6MZZ1_HUMAN
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01-MAY-2000, se
07-FEB-2006, er
Latrophilin-2 (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=20225451; PubMed=10760572; DOI=10.1016/S0167-4781(00)00020-8; White G.R.M., Varley J.M., Heighway J.; "Genomic structure and expression profile of LPHH1, a 7TM gene variably expressed in breast cancer cell lines."; Biochim. Biophys. Acta 1491:75-92(2000).
                                                                                                                                                                                                         05-JUL-2004, integrated into UniProtKB/TrEMBL
05-JUL-2004, sequence version 1.
07-FEB-2006, entry version 6.
Hypothetical protein DKF2p686J2076.
Name=DKF2p686J2076;
                                                                                                                                         Mammalia; Eutheria;
                                                                                                                                                               Eukaryota; Metazoa;
                                                                                                                                                                                     Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=99153747; PubMed=10030676; DOI=10.1038/sj.onc.1202487; White G.R.M., Varley J.M., Heighway J.; "Isolation and characterisation of a human homologue of the latrophilin gene from a region of 1p31.1 implicated in breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
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Q9UJ48;
                   TISSUE=Human
                                            NUCLEOTIDE SEQUENCE.
                                                                                   NCBI_TaxID=9606
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2; Conservative
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                                                                                                                                       Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
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4; Mismatches
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Copyright (c) 1993 - 2006 Biocceleration Ltd.

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114.304 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

The present invention relates to a novel monoclonal antibody (I) that specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. (I) is useful for testing an allergen reactivity of an IgE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy peanut and corn allergens. The sample is a biological sample collected from a dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal antibodies recognise epitopes on canine IgE corresponding to amino acid residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the

Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian IgE or treating asthma or anaphylactic shock.

WPI; 2004-593545/57.

Example 6; Page 9; 14pp; English.

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| 2        |          |          | 9 O      |          |          |          |          |          |          |          |           |          |           |          |          |           |          |          |          | 15 8     |   |
| _        |          |          |          |          |          |          |          |          |          |          |           |          |           |          |          |           |          | _        |          |          |   |
| AAU17678 | ADN69607 | ADN69955 | ADN70395 | ADN68418 | ADN69336 | ADN68615 | ADN67646 | ADN67893 | ABP57763 | ADD43993 | AED55089  | AED49042 | ADY55643  | ADH89500 | ADI12489 | AED61860  | ADN71415 | ADN70579 | ADN70693 | ADN71055 |   |
| Aau17678 | Adn69607 | Adn69955 | Adn70395 | Adn68418 | Adn69336 | Adn68615 | Adn67646 | Adn67893 | Abp57763 | Add43993 | Aed55089  | Aed49042 | Ady55643  | Adh89500 | Adi12489 | Aed61860  | Adn71415 | Adn70579 | Adn70693 | Adn71055 |   |
| 3 Novel  | 7 Human  | 5 Human  | 5 Human  | 3 Human  | 5 Human  | 5 Humar  | 5 Human  | 3 Human  | 3 Human  | CPG2     | _         |          |           | _        | _        |           | 5 Human  | 9 Human  | Human    | -        |   |
|          | n 273    | _        | _        | _        | _        | -        | -        | -        | n pho    | 77       | Hair-bind | mite     | Hair-bind | Scorpion | Scorpion | Chlorotox | n 273    |          |          | -        |   |

## ALIGNMENTS

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RESULT 1
ADR10603
Hammerberg
                                                                                                                                                                                       05-AUG-2004.
                                                                                                                                                                                                                 Equus caballus.
                                                                                                                                                                                                                                      Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody;
                                                                                                                                                                                                                                                                         21-OCT-2004
                                                                                                                                                                                                                                                                                       ADR10603;
                                                                                                                                                                                                                                                                                                   ADR10603 standard; peptide;
                                                                                                                                                           16-JAN-2003; 2003US-0440472P.
                                                                                                                                                                         15-JAN-2004; 2004WO-US003566.
                                                                                                                                                                                                    WO2004065936-A2
                                                                                                                                                                                                                                                         Horse IgE epitope recognised
                                                                                                                                              (UYNC-) UNIV NORTH CAROLINA STATE
                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                    15
                                                                                                                                                                                                                                                          by monoclonal antibody 5.91, SEQ ID
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The invention relates to an immunogenic composition comprising an CC isolated polypeptide having an amino acid sequence that is at least 80% CC identical to 6 (S1-S6), 15 amino acid peptide sequences derived from CC equine immunoglobulin E (the composition induces production of an CC antibody that specifically binds to equine immunoglobulin (Ig)E), the six CC polypeptides are not explicitly identified in the specifically binds (CC polypeptide at least 80% identified in the specifically binds (CC to a polypeptide at least 80% identical to (S1)-(S6), an antibody that specifically binds (CC particularly binds to equine IgE made by the process of immunising an CC animal with a polypeptide at least 80% identical to (S1)-(S6), making an CC animal with a composition further comprising an isolated polypeptide (the antibody that specifically binds to equine IgE (involving immunising an CC animal with a composition further comprising an isolated polypeptide (the CC amino acid sequence of the polypeptide is at least 80% identical to (S1)-(S6)), and collecting antiserum from the animal) and a kit for detection (G equine IgE in a biological sample comprising the antibody and means (CC antibody is useful for detecting equine IgE protein in a biological (CC sample (serum) which involves contacting the sample with the antibody, thus forming an antigen/antibody complex. The antibody and antigen are imposition of the antibody complex and detecting the presence or imposition of the antigen/antibody complex. The antibody and antigen are incompleted to the antibody and antigen are the imposition of the antibody complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and 3.76 were observed to have good cross-reactivity with the epsilon-chain of IgE from cat and horse, but did not exhibit cross-reactivity with either pig or human epsilon-chains of IgE. The present sequence is the horse IgE 5.91 recognition site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunogenic composition comprising an isolated equine immunoglobulin E polypeptide that induces production of antibodies which specifically bind to equine immunoglobulin E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 8; 14pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-765437/72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gershwin LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (REGC ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-NOV-2001; 2001US-00052788
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2003087314-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Equus caballus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Horse; immunoglobulin E; IgE; heavy chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Horse immunoglobulin E, IgE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADC64568;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   heavy chain immunogenic peptide P4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 74; DH o; -
Pred. No. 2.1e-06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunogen; allergy
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RESULT 3
ADR10601
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                              specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. (I) is useful for testing an allergen reactivity of an IgE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut and corn allergens. The sample is a biological sample collected from a dog, cat or horse. (I) is also useful for detecting mammalian IgE and for reating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal antibodies recognise epitopes on canine IgE corresponding to amino acid residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and cat, horse, pig and human by 5.91 and 3.76 was examined.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     complex can be detected. The complex is detected using a second labelled antibody. The peptides are useful for generating antibodies specific for IgE which can serve as a diagnostic test for allergy. The present sequence is a Horse immunoglobulin E, IgE, heavy chain immunogenic peptide from the early portion of the C4 region.
                               of IgE from cat and either pig or human
                                                                                                                                                                                                                                                                                                                                                         Novel antibody that specifically binds to for testing an allergen reactivity of IgE or treating asthma or anaphylactic shock.
                                                                                                                                                                                                                                                                                     The present invention relates to a novel monoclonal antibody (I) that
                                                                                                                                                                                                                                                                                                                       Example 6; Page 9; 14pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-593545/57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-OCT-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADR10601;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADR10601 standard; peptide; 15 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYNC-) UNIV NORTH CAROLINA STATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-JAN-2004; 2004WO-US003566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-AUG-2004
               at, horse, pig and human by 3.71 and 1.70 real reactivity with the epsilon-chain 1.76 were observed to have good cross-reactivity with if IgE from cat and horse, but did not exhibit cross-reactivity with ither pig or human epsilon-chains of IgE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IgE epitope recognised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    66.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             by monoclonal antibody 5.91,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 49;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 7;
0.074;
                                                                                                                                                                                                                                                                                                                                                                            mammalian IgE epitope, useful sample, detecting mammalian IgE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,</u>
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Query Match Best Local Matches

Similarity 9; Conserv

Conservative

55.4%;

Score 41; DB 8 Pred. No. 2.1; 1; Mismatches

8,

Length 15;

4.

Indels

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Sequence 15

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RNNVLIQTDQQATT

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RESULT 5
ADR10602
ID ADR1
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                                                                                                                                                                                                                                        The present invention relates to a novel monoclonal antibody (I) that gpecifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. (I) is useful for testing an allergen reactivity of an IgE sample. The CC allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut can dorn allergens. The sample is a biological sample collected from a CC dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal cantibodies recognise epitopes on canine IgE corresponding to amino acid cresidues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the CC canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from CC cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and CC 3.76 were observed to have good cross-reactivity with the epsilon-chain cat and horse, but did not exhibit cross-reactivity with the cither pig or human epsilon-chains of IgE. The present sequence is the CC pig IgE 5.91 recognition site.
                                                                                                                                                              Query Match
Best Local (
                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian Igor treating asthma or anaphylactic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sus scrofa.
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                                                                                                                                                                                                            Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 6; Page 9; 14pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-593545/57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hammerberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-JAN-2003; 2003US-0440472P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pig IgE epitope recognised by monoclonal antibody 5.91, SEQ ID 7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADR10607 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYNC-) UNIV NORTH CAROLINA STATE
                                                                                 _
                                                                                                                                               Similarity 7; Conserv
                                                                                 RNDAPVOADRHSTTR 15
                                                                                                           RNNVLIQTDQQATTR 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RNDSPIQTDQYTTT 14
                                                                                                                                                                                                                B
                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                              54.1%;
46.7%;
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                                                                                                                                                              Score 40;
Pred. No.
                                                                                                                                                  Mismatches
                                                                                                                                                           DB 3.3;
                                                                                                                                                                              8
                                                                                                                                               4
                                                                                                                                                                            Length 15
                                                                                                                                                  Indels
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                                                                                                                                               Gaps
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ADR10602 standard; peptide; 15

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RESULT 6
ADD44095
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                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian IgE or treating asthma or anaphylactic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal ant
bacterial enzyme; carboxypeptidase G2; CPG2; non-immunogenic; immunogenic; T-cell epitope; MHC class II binding ligand;
                                             CPG2 peptide #1 able to stimulate human T-cells in vitro
                                                                                                          ADD44095;
                                                                                                                                                                                                                                                                                                                                         Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 6; Page 9; 14pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-593545/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hammerberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Felis catus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-OCT-2004
                                                                            15-JAN-2004
                                                                                                                                      ADD44095 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYNC-) UNIV NORTH CAROLINA STATE
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                                                                                                                                                                                                                                                                               Score 37; DB
Pred. No. 11;
4; Mismatches
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Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        carboxypeptidase G2 (CPG2). The modified enzyme can result in CPG2 proteins that are substantially non-immunogenic or less immunogenic than any non-modified CPG2 having essentially the same biological specificity when used in vivo, and comprising specific amino acid residues having alterations compared with the non-modified parochial enzyme. The alterations cause a reduction or an elimination of one or more of T-cell epitope sequences, which act in the parental enzyme as MHC class II binding ligands and stimulate T-cells. The modified CPG2 enzyme and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-NOV-2001;
25-JAN-2002;
13-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CPG2 proteins have immunostimulant activity and may be used in enzyme therapy. The modified CPG2 enzyme may be used to induce an immune response in a human host, or as a therapeutic entity such as the gene directed enzyme prodrug strategy. The peptide is useful for the manufacture of a modified CPG2 enzyme having substantially no or less immunogenicity than any non-modified parental enzyme when used in vivo, and for vaccination of patients to reduce immunogenicity to CPG2 in vivo.

This sequence represents a CPG2 enzyme peptide able to stimulate human T-
                               bacterial enzyme; carboxypeptidase G2; CPC2; non-immunogenic; immunogenic; T-cell epitope; MHC class II binding ligand; immunostimulant; enzyme therapy; immune response; gene directed enzyme prodrug strategy; vaccine; enzyme; EC 3.4.17.11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New modified bacterial enzyme carboxypeptidase G2 (CPG2) having substantially non-immunogenic or less immunogenic than any non-modified CPG2, useful for inducing an immune response in a human host.
Pseudomonas sp. RS-16
                                                                                                                         Carboxypeptidase G2 (CPG2) enzyme immunogenic peptide #1.
                                                                                                                                                              15-JAN-2004
                                                                                                                                                                                               ADD44240;
                                                                                                                                                                                                                                ADD44240 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel modified bacterial enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3; Page 14; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hellendoorn K,
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gene directed enzyme prodrug strategy; vaccine; enzyme; EC 3.4.17.11.
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                     RNNVLIQ--TDQQ 11
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2002EP-00001778.
2002EP-00020634.
                                                                                                                                                            (first
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                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                           43.2%;
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Pred. No. 94;
2; Mismatches
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CC proteins that are substantially non-immunogenic or less immunogenic than
CC any non-modified CPG2 having essentially the same biological specificity
CC when used in vivo, and comprising specific amino acid residues having
CC alterations compared with the non-modified parochial enzyme. The
CC alterations cause a reduction or an elimination of one or more of T-cell
CC pitope sequences, which act in the parental enzyme as MHC class II
CC binding ligands and stimulate T-cells. The modified CPG2 enzyme and the
CC CPG2 proteins have immunostimulant activity and may be used in enzyme
CC therapy. The modified CPG2 enzyme may be used to induce an immune
CC tresponse in a human host, or as a therapeutic entity such as the gene
CC directed enzyme prodrug strategy. The peptide is useful for the
CC manufacture of a modified CPG2 enzyme having substantially no or less
CC immunogenicity than any non-modified parental enzyme when used in vivo,
CC and for vaccination of patients to reduce immunogenicity to CPG2 in vivo.
CC This sequence represents an immunogenic peptide of the CPG2 enzyme of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local
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25-JAN-2002;
13-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New modified bacterial enzyme carboxypeptidase G2 (CPG2) having substantially non-immunogenic or less immunogenic than any non-modified CPG2, useful for inducing an immune response in a human host.
                                                                                                                               US2005181388-A1
                                                                                                                                                                                                                                                                                      protein purification; antibacterial; antimicrobial; infection;
                                                                                                                                                                                                                                                                                                                                                    Pseudomonas aeruginosa phosphoglucosamine mutase peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AEC11192 standard; peptide; 12
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                                                                                                                                                                                                                                                     screening; phosphoglucosamine mutase
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2002EP-00001778.
2002EP-00020634.
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04-OCT-2004; 2004US-00958216

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02-APR-2002
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2002US-0436987P.
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2002US-0437013P.
2002US-0437038P.
2002US-0437281P.
2002US-0437527P.
2002US-0437527P.
2002US-0437620P.
2002US-0437638P.
2003WO-CA000464.
2003WO-CA000464.
2003WO-CA000465.
 Dharamsi A,
Buzadzija K,
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2002US-038639P
2002US-038680P
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2002US-0424033P
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2002US-0425200P
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2002US-0370792P.
2002US-0370829P.
2002US-0370829P.
2002US-0370899P.
2002US-0371067P.
2002US-0371107P.
2002US-03711185P.
2002US-0371185P.
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Arrowsmith C,
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Ouyang H, Pind
Virag C;
                                                                                                                                                                  23-APR-1998;
09-SEP-1998;
08-JAN-1999;
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Paige LA,
Mcdonnell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oestrogen receptor; estrogen; estradiol; oestrogen response element; ERE; binding; biological activity; fingerprint; molecular braille; cellular braille; modulation; tamoxifen; breast cancer; ovarian cancer; menopause; osteoporosis; selective oestrogen receptor modulator; identification; characterisation; classification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
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                                                                                                                                                                                                                                                                                                            26-MAR-1999;
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                                   Hamilton PT,
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                                                                                                                                                                  98US-0082756P.
98US-0099656P.
99US-0115345P.
   Christensen
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                                                                                                  PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide;
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                                   Fowlkes
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Pred. No. 2.6e
6; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mcdonald M, Nethery-Brokx K,
Tai M, Thalakada R, Vallee F;
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                                   Buehrer B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.6e+02;
2;
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                                      Barnett T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                is used
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WPI; 2000-013281/01

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a method for predicting the biological CC activity of new receptor modulating compounds (I) using novel oligomeric CC peptides (biokeys) which have differential abilities to bind to 2 conditions. The method is used to identify new CC drugs that are physiological or pharmacological agonists/antagonists and CC that target various receptors, which are involved in certain disease CC conditions. The system may be used as a primary screening tool to CC identify hits, to classify lead compounds from a drug screen to, CC characterise selective oestrogen receptor modulators (SERMs) in terms of CC agonist and antagonist function and to predict possible clinical effects of SERMs such as tissue and receptor specificity. The method can also be capplied to the fractionation of mixtures of SERMs to determine which CC applied to the fractionation of mixtures of SERMs to determine which CC applied to the fractionity of the prospector, to identify, characterise and classify CC epidermal growth factor, to identify, characterise and classify CC epidermal growth factor, to identify, characterise and classify CC used to modulators of receptor activity. Peptides comprising a LXXLL motif may be contained and ameliorating the effects of menopause, including contention and oestrogen receptor in treating e.g. breast and costeoporosis. AAY65439 to AAY65652 represent oestrogens receptor, cestradiol receptor and oestrogen response element binding peptides given in the exemplification of the present invention. AAZ35740 to AAZ35745
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Best Local (
                                 WPI; 2002-154969/20
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21-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oestrogen receptor; breast cancer; combinatorial peptide library;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oestrogen receptor beta binding peptide 17E-beta.
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                                                                                             Fowlkes DM,
                                                                                                                                                          (KARO-) KARO
                                                                                                                                                                                                                                                                                                                        11-JUL-2001; 2001WO-US021867
                                                                                                                                                                                                                                                                                                                                                                                           17-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200204956-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   receptor modulating compound.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                             Barnett TR,
                                                                                                                                                          BIO USA INC
                                                                                                                                                                                                                      2000US-00614865
2001US-00860688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                             Buehrer
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Identifying receptor-binding peptides comprises screening combinatorial peptide library presented in form of cells each of which coexpress one peptide member and receptor with signal producing system for reporting
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Disclosure; Page 146; 175pp; English

Comprising screening a combinatorial library of peptides, comprising screening a combinatorial peptide library presented in the compress the receptor or its ligand-binding receptor comoiety and one member of the library, together with a signal producing system for reporting binding of the peptide to the receptor. Also compound which modulates the biological activity of a receptor activity of a compound by (i) product the receptor by the method above, (b) consider the members differ in their ability to bind the receptor. Move the receptor by the method above, (c) in identifying peptides which bind the receptor is in, where the field peptides, compound by (i) providing a panel of identified peptides, compound by (i) providing a panel of identified peptides, compound by (i) providing a panel of identified peptides, coff a number of treference substances known to modulate the biological activity of a compound by (i) providing a panel of identified peptides, coff a number of treference substances known to modulate the biological activity of the receptor on the binding of each member of the panel is comparised as a reference fingerprint for each reference coffice the panel to the receptor, thereby obtaining a test finderprint for the test substance, (ii) screening a test substance of unknown activity relative the panel to the receptor, thereby obtaining a test finderprint for the test substance and (iv) predicting the biological activity of the test compound (e.g. and (iv) predicting the biological activity of the test substance with similar finderprint. The members are useful for identifying a binding peptide which binds a cutivity of a compound (e.g. antiogen) receptor. The identified peptides which binds to the receptor activity of a compound (e.g. antiogen) receptor. The identified peptides which is different conceptors, the superior can be applied to a variety of small organic method has several advantages over whole animal-based assay systems in that the same technology can be applied to a variety of different Sequence 15 AA; The invention relates to identifying a binding peptide which binds a receptor and which is a member of a combinatorial library of peptides

밁 Ş Matches Query Match Best Local ω 3 NVLIQTDQQATTR 15 6 Similarity NLLCLLDQEACSR 15 Conservative 39.2%; Score 29; DB 5; Pred. No. 3.3e+02; 3; Mismatches 4 DB 5; Length 15; 4. ٠<u>.</u> Gaps

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ADM79070 standard; peptide; 15

Oestrogen receptor beta binding peptide #57 03-JUN-2004 (first entry

binding peptide; drug indentificaton; in vivo imaging; oestrogen receptor beta.

Unidentified

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TAREST CONTRACTOR OF CONTRACTO
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21-MAY-2001;
11-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a method of identifying a binding peptide which binds a receptor comprises screening a combinatorial peptide library presented in cells which co-expresses one member peptide and the receptor, together with a signal producing system for reporting binding. The method is useful in identifying drugs which can mediate the biological activity of a target protein. The receptor-binding library members may be used as therapeutic or diagnostic reagents, for in vivo imaging, or to purify target from a fluid, e.g. blood. The present sequence represents an oestrogen receptor beta binding peptide.
WPI; 2004-593545/57.
                                                                                                                                                                                                                                                                           05-AUG-2004
                                                                                                                                                                                                                                                                                                                              WO2004065936-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sheep IgE epitope recognised by monoclonal antibody 5.91,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-OCT-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and a signal producing system.
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NLLCLLDQEACSR 15
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2001US-00860688.
2001WO-US021867.
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3.3e+02;
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XFFFX8X33333333333333333333333333333
                                                                                                                                                            Novel antibody that specifically binds to mammalian IgE epitope, usefu for testing an allergen reactivity of IgE sample, detecting mammalian or treating asthma or anaphylactic shock.
                                                                                                                                                                     IgE
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Page 9; 14pp; English.

specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. (I) is useful for testing an allergen reactivity of an IgE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut and corn allergens. The sample is a biological sample collected from a dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal antibodies recognise epitopes on canine IgE corresponding to amino acid residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from cat, horse, pig and human by 5.91 and 3.76 were observed to have good cross-reactivity with the epsilon-chain of IgE from cat and horse, but did not exhibit cross-reactivity with either pig or human epsilon-chains of IgE. The present sequence is the Sequence 15 AA; The present invention relates to a novel monoclonal antibody (I) IgE 5.91 recognition site.

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Matches
                                       Query Match
Best Local
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                \vdash
                               Similarity 6; Conserv
              RNNVLIQTDQQATTR 15
RNKELMREGOHTTTO 15
                                Conservative
                                       39.2%;
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                                        Score 29;
Pred. No.
                                Mismatches
                     ; DB 8;
5. 3.3e+02;
6; Indels
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В á

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RESULT 13
AEC98485
ID AEC98
XX
AC CONTRACTOR OF A CONTRACTOR 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AEC98485 standard; peptide; 15
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(INNO-) (CHES/) (NEWM/) (NOTH/) (BAKE/) (SOUT/) (BABE/) (CHEN/) (CHEN/) (HUAN/)

SOUTHWOOD BABE L M. CHEN Y. MOTHE B. BAKER D.

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DEYOUNG L M. HUANG M T F. POWER S D.

31-DEC-2003; 02-JUL-2004; 03-JAN-2005;

2003US-0533211P. 2004US-0584652P. 2005WO-US000077.

EPIM-)

EPIMMUNE INC

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INNOGENETICS I CHESNUT R. NEWMAN M J.

WO2005089164-A2 Human papillomavirus.

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RESULT 14
AEC98272
ID AEC98
XX AEC98
AC AEC96
AC AEC91
AC AEC96
AC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC The invention relates to a new polynucleotide comprising a multi-epitope CC construct comprising nucleic acids encoding the human papillomavirus CC (HPV) cytotoxic Tlymphocyte (CTL) epitopes e.g., HPV16.E1.214, and that CC are directly or indirectly joined to one another in the same reading CC frame, a vaccine minigene. Also included are a vector comprising the CC multi-epitope construct, a polypeptide comprising an amino acid sequence cc encoded by the polynucleotide, a composition (comprising the polynucleotide, vector or polypeptide and a carrier), a cell CC polynucleotide, vector polypeptide and a carrier), a cell CC immune response against human papillomavirus virus (HPV) and making the polynucleotide, vector or polypeptide), inducing an CC different strains of HPV and are from the El, E2, E6 and E7 proteins. The epitopes may be linked via a GP-anchor/spacer peptide. The order of the cpitopes may be linked via a GP-anchor/spacer peptide. The order of the cpitopes in the vaccine protein are disclosed in the tables referred to cin the claims of the specification. The polynucleotide, vector or cpolypeptide is useful in preparing a composition for inducing an immune CC defense against HPV infection and HPV-related cancers. The present contents of the sequence is an HPV-derived peptide epitope that binds to a subclass of the human leukocyte antigen (HLA) peptides.
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Best Local S
Matches 4
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Chen Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human papilloma virus infection; virucide; cancer; cytostatic; vaccine, epitope mapping; immune stimulation; cytotoxic T-lymphocyte; human leukocyte antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HLA-DR binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AEC98272 standard; peptide;
                                                                                                                                                                           31-DEC-2003;
02-JUL-2004;
                                                                                                                                                                                                                                                           03-JAN-2005; 2005WO-US000077.
                                                                                                                                                                                                                                                                                                                    29-SEP-2005
                                                                                                                                                                                                                                                                                                                                                                         WO2005089164-A2
                                                                                                                                                                                                                                                                                                                                                                                                                           Human papillomavirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AEC98272;
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                                   NEWM/)
                                                           CHES/)
                                                                                        INNO-)
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                                                        INNOGENETICS NV. CHESNUT R.
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Deyoung LM,
                                                                                                                   EPIMMUNE INC.
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     MOTHE B.
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                                                                                                                                                                           2003US-0533211P.
2004US-0584652P.
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40.0%;
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duang MTF,
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Pred. No.
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Power SD;
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                                                                                                                        CC construct comprising nucleic acids encoding the human papillomavirus (HPV) cytotoxic Tlymphocyte (CTL) epitopes e.g., HPV16.El 214, and that CC are directly or indirectly joined to one another in the same reading CC frame, a vaccine minigene. Also included are a vector comprising the CC multi-epitope construct, a polypeptide comprising an amino acid sequence CC encoded by the polynucleotide, a composition (comprising the polynucleotide, vector or polypeptide and a carrier), a cell CC (comprising the polynucleotide, vector or polypeptide and a carrier) a cell CC (immune response against human papillomavirus virus (HPV) and making the CC polynucleotide, vector or polypeptide. The epitopes are derived from CC different strains of HPV and are from the El, E2, E6 and E7 proteins. The CC epitopes may be linked via a GP-anchor/spacer peptide. The order of the CC epitopes in the vaccine protein are disclosed in the tables referred to CC polypeptide is useful in preparing a composition for inducing an immune CC response against HPV infection and HPV-related cancers. The present CC sequence is an HPV-derived peptide epitope that binds to a subclass of CC human leukocyte antigen (HLA) peptides.
                                                          Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chesnut
Chen Y,
                                                                                                                       Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (BAKE/)
(SOUT/)
(BABE/)
(CHEN/)
(CHEN/)
(DEYO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 315;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New polynucleotide comprises a multi-epitope construct comprising nucleic acids encoding the human papillomavirus (HPV) cytotoxic T lymphocyte (CTL) epitopes, useful in preparing a vaccine against HPV.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R, No.
\vdash
                             5 LIQTDQQATT 14
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CHEN Y.
                                                            4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HUANG M T POWER S D.
                                                                           Similarity
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MVQVEEQQTT
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                                                            Conservative
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                                                                         37.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    518pp; English.
                                                          Score 28; DB 9;
Pred. No. 5e+02;
1; Mismatches
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Power
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       comprising a
                                                            <u>ب</u>
                                                                                        Length 15;
                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     multi-epitope
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                                                            Gaps
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Query Match
Best Local Similarity 40.0%; Pred. No. 5e+02;
Matches 4; Conservative 4; Mismatches 2; Indels 0; Gaps 0;

Qy 5 LiqTDQQATT 14

:: | : | | |
Db 1 MVQVEEQQTT 10

RESULT 15
AAG95534
ID AAG95534;
ID AAG95534;
XX
AAG95534
ID 18-SEP-2001 (first entry)
XX
DE Human complementary peptide; 10 AA.
XX
Human complementary peptide; ligand; drug discovery; drug design.
XX
OS Homo sapiens.
XX
PN WO200142277-A2.
XX
PN WO200142277-A2.
PF 13-DEC-2000; 2000WO-GB004776.
XX
PR 13-DEC-1999; 99GB-00029464.
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Search completed: May 30, 2006, 10:12:42 Job time: 62 secs
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                                                                                                                                                                                                                        SO CCCCCXXX PT PT XXX PA
                                                                                                                                            Query Match 36.5%;
Best Local Similarity 71.4%;
Matches 5; Conservative
                                                                                                                                                                                                                                                    The invention relates to a set of complementary peptide ligands generated from the human genome. The complementary peptides interact with their relevant target proteins encoded in the human genome. They can be used as reagents in drug discovery and as lead ligands to facilitate drug design and development. The present sequence is a complementary peptide provided in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                  A set of peptide ligands consisting of specific complementary peptides to proteins encoded by genes of the human genome, useful in an assay for screening and identifying of one or more novel peptides which are drug candidates or pro-drugs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-408419/43.
                                                                                                                                                                                                                      Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                              Example 4; Page 290; 646pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Roberts GW, Heal JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PROT-) PROTEOM LTD.
                                                                       1 SDQQTTT 7
                                                                                                           8 TDQQATT 14
                                                                                                                                            Score 27; DB 4; Length 10; Pred. No. 4.9e+02; 1; Mismatches 1; Indels
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Result
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Maximum Match 100%
Listing first 45 summaries
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Sequence:
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BB
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  May 30, 2006, 10:21:55; Search time 47 Seconds (without alignments)
147.835 Million cell updates/sec
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Match Length
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74
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| / EMC_Celerra_SIDS3/ptodate3/2/pubpaa/US08_PUBCOMB.pep:*
| / EMC_Celerra_SIDS3/ptodate3/2/pubpaa/US08_PUBCOMB.pep:*
| / EMC_Celerra_SIDS3/ptodate3/2/pubpaa/US09_PUBCOMB.pep:*
| / EMC_Celerra_SIDS3/ptodate3/2/pubpaa/US10A_PUBCOMB.pep:*
| / EMC_Celerra_SIDS3/ptodate3/2/pubpaa/US10B_PUBCOMB.pep:*
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                                      US-10-758-165-3

US-10-052-788-4

US-10-758-165-1

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US-10-958-216-713

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US-10-958-216-713

US-10-958-216-713

US-10-958-216-713

US-10-758-165-4

US-09-9572-4048-1728

US-10-497-091-129

US-10-497-091-129

US-10-497-091-129

US-10-497-091-129

US-10-497-091-123

US-10-497-091-123

US-10-497-091-123

US-10-21-125-1

US-10-225-567A-1123

US-11-004-270-94

US-11-004-273-94

US-11-004-273-94

US-11-004-273-560

US-10-895-064-2404
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                  APPLICANT: Gershwin, Laurel J.

APPLICANT: Hettigrew, Howard David
APPLICANT: Kalina, Warren V.

APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Epsilon Immunoglobulin Chain Derived Peptides for
TITLE OF INVENTION: Induction of Anti-IgE Antibodies
FILE REFERENCE: 023070-121000US
FURRENT APPLICATION NUMBER: US/10/052,788
CURRENT FILING DATE: 2001-11-08
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
LENGTH: 15
TYPE: PRT
ORGANIGM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; LENGTH: 15
; TYPE: PRT
; ORGANISM: Equus caballus
US-10-758-165-3
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US-10-052-788-4
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                                                                                                                                                                                                                                                                                                                                                 Sequence 4, Application US/10052788 Publication No. US20030087314A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 15; Conserva
      FEATURE:
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| 24 32.4 9 6 US-11-129-741-2404 24 32.4 10 4 US-10-117-937-561 24 32.4 10 6 US-11-067-064-561 24 32.4 10 6 US-11-067-159-561 24 32.4 12 4 US-10-097-175-66 24 32.4 12 4 US-10-264-309-289 24 32.4 12 4 US-10-264-309-289 24 32.4 13 3 US-09-791-377-584 24 32.4 13 3 US-09-791-377-584 24 32.4 13 4 US-10-014-340-569 24 32.4 13 4 US-10-014-340-569 24 32.4 13 4 US-10-014-38-7 24 32.4 13 4 US-10-369-736-31 24 32.4 13 4 US-10-369-736-31 24 32.4 13 4 US-10-369-738-31 24 32.4 13 4 US-10-369-738-31 24 32.4 13 5 US-10-808-187-706 24 32.4 13 5 US-10-807-807-706 24 32.4 13 6 US-11-028-058-31  | 32.4 10 6 US 32.4 10 6 US 32.4 10 6 US 32.4 12 4 US 32.4 12 4 US 32.4 12 3 US 32.4 13 3 US 32.4 13 3 US 32.4 13 4 US 32.4 13 6 US   | 45               | 44                | 43                | 42               | 41 | 40 | 39 | 38 | 37 | 36 | 35                | 34                | 33                | 32               | 31                | 30                | 29                | 28                 |
|--|---|------------------|-------------------|-------------------|------------------|----|----|----|----|----|----|-------------------|-------------------|-------------------|------------------|-------------------|-------------------|-------------------|--------------------|
| .4.4.110 6 US .4.4.112 4 US .4.4.113 3 US .4.4.113 4 US .4 | .4 9 6 US-11-129-741-2404 Sequence .4 10 4 US-10-117-337-561 Sequence .4 10 6 US-11-067-064-561 Sequence .4 10 6 US-11-067-0561-561 Sequence .4 11 6 US-11-0697-175-66 Sequence .4 12 4 US-10-097-175-66 Sequence .4 12 4 US-10-264-309-289 Sequence .4 12 5 US-10-264-309-289 Sequence .4 13 3 US-09-791-377-584 Sequence .4 13 4 US-10-014-340-568 Sequence .4 13 4 US-10-014-340-569 Sequence .4 13 4 US-10-014-338-7 Sequence .4 13 4 US-10-014-338-7 Sequence .4 13 4 US-10-13-582-14 Sequence .4 13 4 US-10-13-582-14 Sequence .4 13 5 US-10-807-808-31 Sequence .4 13 5 US-10-807-808-31 Sequence .4 13 5 US-10-807-808-31 Sequence .4 13 6 US-11-028-058-31 Sequence  | 24               | 24                | 24                | 24               | 24 | 24 | 24 | 24 | 24 | 24 | 24                | 24                | 24                | 24               | 24                | 24                | 24                | 24                 |
| 8 20 20 20 20 20 20 20 20 20 20 20 20 20   | 6 US-11-129-741-2404 4 US-10-117-937-561 5 Equence 6 US-11-067-564-561 5 Equence 6 US-11-067-566 6 US-11-067-159-561 6 US-11-067-159-561 6 US-10-097-175-66 6 US-10-264-309-289 5 US-10-264-309-289 5 US-10-264-309-289 6 US-10-14-340-568 6 US-10-14-340-568 6 US-10-014-340-569 6 US-10-014-340-569 6 US-10-014-340-569 6 US-10-014-340-569 6 US-10-112-582-14 6 US-10-112-582-14 6 US-10-807-807-706 7 US-10-807-807-706 8 Equence 7 US-10-807-807-706 8 Equence 8 US-11-0807-807-706 8 Equence 9 US-11-0807-807-706 8 Equence 9 US-11-0807-807-706 8 Equence 9 US-11-0807-807-807-706 8 Equence 9 US-11-0807-807-807-706 8 Equence 9 US-11-0807-807-807-807-807-807-807-807-807-80  | 32.4             |                   |                   |                  |    |    |    |    |    |    |                   |                   |                   |                  |                   |                   |                   |                    |
|  | US-10-117-937-561 Sequence US-10-117-937-561 Sequence US-11-067-064-561 Sequence US-11-067-159-561 Sequence US-11-067-159-561 Sequence US-10-264-309-289 Sequence US-10-264-309-289 Sequence US-09-791-378-584 Sequence US-09-791-377-584 Sequence US-09-791-377-584 Sequence US-09-791-377-589 Sequence US-09-791-370-569 Sequence US-09-791-370-569 Sequence US-10-014-338-7 US-10-014-338-7 US-10-014-338-7 US-10-018-582-14 Sequence US-10-112-582-14 Sequence US-10-112-582-14 Sequence US-10-1808-187-706 Sequence US-10-807-807-706 Sequence US-11-028-058-31 Sequence   | 13               | 13                | 13                | 13               | 13 | 13 | 13 | 13 | 13 | 13 | 13                | 12                | 12                | 12               | 10                | 10                | 10                | v                  |
| US-11-129-741-2404 US-11-10-17-937-561 US-11-067-159-561 US-11-067-159-561 US-11-0-264-309-289 US-10-264-309-289 US-10-264-309-289 US-10-264-309-289 US-10-014-340-568 US-10-014-340-569 US-10-014-340-569 US-10-14-338-7 US-10-14-338-7 US-10-1582-14 US-10-1808-187-706 US-10-807-807-706 US-11-807-807-706 US-11-808-087-31   | 11-129-741-2404 Sequence -10-117-937-561 Sequence -11-067-159-561 Sequence -10-097-175-66 Sequence -10-264-309-289 Sequence -10-264-309-289 Sequence -10-2791-377-584 Sequence -09-791-377-584 Sequence -10-014-340-568 Sequence -10-014-340-569 Sequence -10-014-340-569 Sequence -10-014-340-706 Sequence -10-014-38-7 Sequence | σ                | رى<br>د           | ഗ                 | 4                | 4  | 4  | 4. | 4  | 4. | w  | W                 | ŋ                 | 4                 | 4.               | O                 | σ                 | 4                 | σ                  |
|  |   | US-11-028-058-31 | US-10-807-807-706 | US-10-808-187-706 | US-10-369-738-31 |    |    |    |    |    |    | US-09-791-378-584 | US-10-264-309-289 | US-10-264-309-289 | US-10-097-175-66 | US-11-067-159-561 | US-11-067-064-561 | US-10-117-937-561 | US-11-129-741-2404 |

ALIGNMENTS

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Sequence 3, Application US/10758165

Publication No. US20050196816A1

GENERAL INFORMATION:

APPLICANT: Hammerberg, Bruce
ITILE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN

FILE REFERENCE: 5051-661

CURRENT APPLICATION NUMBER: US/10/758,165

CURRENT FILING DATE: 2004-01-16

PRIOR APPLICATION NUMBER: US/10/758,165

PRIOR APPLICATION NUMBER: US/10/758,165

PRIOR PILING DATE: 2003-01-16

NUMBER OF SEQ ID NOS: 16

NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                             SOFTWARE: PatentIn version 3.2 SEQ ID NO 3
1 RNNVLIQTDQQATTR 15
                      1 RNNVLIQTDQQATTR 15
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                                                                                  Score 74; DB 5;
Pred. No. 2.5e-06;
Mismatches 0;
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                                                                                                                               Length 15;
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CURRENT APPLICATION NUMBER: US/10/758,165
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: US 60/440,472
; PRIOR FILING DATE: 2003-01-16
; NUMBER OF SEO ID NOS: 16
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 1
; SEQ ID NO 1
; LENGTH: 15
; TYPE: PRT
; ORGANISM: Canis familiaris
US-10-758-165-1
                                                                                                                                ; LENGTH: 15
; TYPE: PRT
; ORGANISM: Sus s
US-10-758-165-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Description of Artificial Sequence:epitope peptide
; OTHER INFORMATION: P4, early portion of C4 of equine IgE epsilon
; OTHER INFORMATION: heavy chain
US-10-052-788-4
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US-10-758-165-7
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                                                               Query Match
Best Local S
Matches 7
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Best Local Similarity 64.3
Conservative
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TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR FILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn version 3.2
SEQ ID NO 7
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Publication No. US20050196816A1
GENERAL INFORMATION:
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Publication No. US20050196816A1
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TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661
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                            RNNVLIQTDQQATTR 15
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f; Pred. No. 0.0
0; Mismatches
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Pred. No. 2.2;
1; Mismatches
                                                             Score 40; DB 5;
Pred. No. 3.3;
4; Mismatches
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RESULT 5

Sequence 713, Application US/10958216
Publication No. US20050181388A1
GENERAL INFORMATION:
APPLICANT: EDWARDS, ALED
APPLICANT: DHARAMSI, AKIL

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; LENGTH: 15
; TYPE: PRT
; ORGANISM: Felis catus
US-10-758-165-2
RESULT 7
US-10-958-216-713
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                                                                                                                                                                                                                            ; ORGANISM: homo sapiens US-10-497-091-128
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Publication No. US20050196816A1

GENERAL INFORMATION:

APPLICANT: Hammerberg, Bruce
TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661

CURRENT APPLICATION NUMBER: US/10/758,165

CURRENT FILING DATE: 2004-01-16

PRIOR APPLICATION NUMBER: US 60/440,472

PRIOR FILING DATE: 2003-01-16

NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 312
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 128
LENGTH: 15
                                                                                                                                                 Query Match 43.3
Best Local Similarity 61.9
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn version 3.2 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 128, Application US/10497091 Publication No. US20050074863A1
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Best Local
                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: EP01128519.4 PRIOR FILING DATE: 2001-11-29
                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: PCT/EP02/13351
PRIOR FILING DATE: 2002-11-27
PRIOR APPLICATION NUMBER: EP02020634.8
PRIOR FILING DATE: 2002-09-13
PRIOR APPLICATION NUMBER: EP02001778.6
PRIOR FILING DATE: 2002-01-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: T-CELL EPITOPES IN CARBOXYPEPTIDASE FILE REFERENCE: MER-130 CURRENT APPLICATION NUMBER: US/10/497,091 CURRENT FILING DATE: 2004-05-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: HELLENDOORN, Koen APPLICANT: BAKER, Matthew APPLICANT: WILLIAMS, Steven
                                                                                                                                                                                                                                                                      TYPE: PRT
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                                                                            RDNVLFQAATDEQ 15
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Pred. No. 11;
4; Mismatches
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APPLICANT: APPLICANT:

VEDADI,

MASOUD

ALAM, MUHAMMAD ZAHOOR

ARROWSMITH,

, DONALD E.

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RESULT 8
US-10-346-162-192
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                                                                                                                                                                                                                                                                                                                                                               ; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-10-958-216-713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/369,511
PRIOR FILING DATE: 2002-04-02
PRIOR APPLICATION NUMBER: 60/385,089
PRIOR FILING DATE: 2002-05-31
PRIOR APPLICATION NUMBER: 60/385,751
PRIOR APPLICATION NUMBER: 60/386,553
                                                                         Sequence 192, Application US/10346162 Publication No. US20030224390A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                         Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 713
                                                                                                                                                                                                                                                                                       Matches
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CURRENT APPLICATION NUMBER: US/10/958,216
CURRENT FILING DATE: 2004-10-04
PRIOR APPLICATION NUMBER: PCT/CA03/00462
PRIOR FILING DATE: 2003-04-02
PRIOR FILING DATE: 2003-04-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/386,390
PRIOR FILING DATE: 2002-06-06
PRIOR APPLICATION NUMBER: 60/386,601
PRIOR FILING DATE: 2002-06-06
                                       APPLICANT: KARO BIO USA, INC. APPLICANT: FOWLKES, Dana M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: VIRAG, CRISTINA
TITLE OF INVENTION: NOVEL PURIFIED POLYPEPTIDES FROM BACTERIA
FILE REFERENCE: IPT-205.01
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                                                                                                                                                                                                                                                                                                       Match 39.2%;
Local Similarity 33.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/386,367
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                   FOWLKES, BARNETT,
BARNETT, Thomas R. BUEHRER, Benjamin
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VALLEE, FRANCOIS
VIRAG, CRISTINA
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MCDONALD, MERRY-LYNN
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CANADIEN, VERONICA
DOMAGALA, MEGAN
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PINDER, BENJAMIN
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Pred. No. 2.4e+02;
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RESULT 10
US-09-572-404B-1728
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US-10-758-165-4
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; OTHER INFORMATION: synthetic
US-10-346-162-192
                                                 Sequence 1728, Application US/09572404B
Publication No. US20030078374A1
GENERAL INFORMATION:
APPLICANT: Protecom Ltd
TITLE OF INVENTION: Complementary peptide ligands from the human genome
FILE REFERENCE: Human patent
CURRENT APPLICATION NUMBER: US/09/572,404B
CURRENT FILING DATE: 2000-05-17
NUMBER OF SEQ ID NOS: 4203
SOFTWARE: ProtePatent version 1.0
SEQ ID NO 1728
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 4
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Publication No. US20050196816A1
GENERAL INFORMATION:
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SEQ ID NO 192
LENGTH: 15
TYPE: PRT
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR FILING DATE: 2003-01-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Hammerberg, Bruce
TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661
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ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
                                      ENGTH: 10
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Local Similarity 40.0%;
les 6; Conservation
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Local Similarity 46.2%;
1es 6; Conservative
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 Mismatches

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Pred. No. 3.1e+02;
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Pred. No. 3.1e+02;
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RESULT 12
US-10-172-425B-37
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; ORGANISM: homo sapiens
US-10-497-091-14
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Best Local Similarity
Thes 7; Conserva
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Publication No. US20030147908A1
GENERAL INFORMATION:
APPLICANT: Kaempfer, Raymond
APPLICANT: Arad, Gila
               SEQ ID
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LENGTH: 13
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                                                                                                        TITLE OF INVENTION: BROAD SPECTRUM ANTAGONISTS AND VACCINES TITLE OF INVENTION: DIRECTED ACAINST PYCOCRNIC EXOTOXINS FILE REBERENCE: A31967-PCT-USA-A-A 06031.0164 CURRENT APPLICATION NUMBER: US/10/172,425B CURRENT FILING DATE: 2002-06-13 PRIOR APPLICATION NUMBER: 09/150,947 PRIOR FILING DATE: 1998-09-10 PRIOR APPLICATION NUMBER: PCT/IL97/00438 PRIOR APPLICATION NUMBER: PCT/IL97/00438 PRIOR APPLICATION NUMBER: ISRAEL 119938 PRIOR FILING DATE: 1997-12-30 PRIOR FILING DATE: 1996-12-30 PRIOR PRIOR FILING DATE: 1996-12-30 PRIOR FILING DATE: 1996-12-30 PRIOR FILING DATE: 1996-12-30 PRIOR FILING DATE: 1996-12-30 PRIOR PRIOR FILING DATE: 1996-12-30 PRIOR FILING DATE: 1996-12-30 PRIOR PRIOR FILING DATE: 1996-12-30 PRIOR PRIOR PRIOR FILING DATE: 1996-12-30 PRIOR PRIOR
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NUMBER OF SEQ ID NOS: 57
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 37
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PRIOR FILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: EP01128519.4
PRIOR FILING DATE: 2001-11-29
NUMBER OF SEQ ID NOS: 312
SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR FILING DATE: 2002-11-27
PRIOR APPLICATION NUMBER: EP02020634.8
PRIOR FILING DATE: 2002-09-13
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TITLE OF INVENTION: T-CE
FILE REFERENCE: MER-130
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APPLICANT: BAKER, Matthew
APPLICANT: WILLIAMS, Steven
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OTHER INFORMATION:
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No. US20050074863A1
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58.3%;
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Pred. No. 4.5e+02;
1; Mismatches 1
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Pred. No. 6.1e+02
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APPLICANT: Huang, Xueying
APPLICANT: Wang, Hong
APPLICANT: Wu, Ying
TITLE OF INVENTION: Peptide-Based Conditioners and Col
TITLE OF INVENTION: Nails
FILLE REFERENCE: CL2296 US NA
CURRENT APPLICATION NUMBER: US/10/935,642
CURRENT FILING DATE: 2004-09-07
NUMBER OF SEQ ID NOS: 104
SOFTWARE: Patentin version 3.2
SEQ ID NO 48
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US-10-935-642-48
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; TYPE: PRT
; ORGANISM: homo sapiens
US-10-497-091-129
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US-10-497-091-129
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CURRENT APPLICATION NUMBER: US/10/497,091
CURRENT FILING DATE: 2004-05-28
PRIOR APPLICATION NUMBER: PCT/EP02/13351
PRIOR FILING DATE: 2002-11-27
PRIOR APPLICATION NUMBER: EP02020634.8
PRIOR FILING DATE: 2002-09-13
PRIOR APPLICATION NUMBER: EP02001778.6
PRIOR APPLICATION NUMBER: EP02001778.6
PRIOR FILING DATE: 2002-01-25
PRIOR FILING DATE: 2002-01-25
                                                                                                                                                                                                                                                                                                                        Sequence 48, Application US/10935642 Publication No. US20050050656A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 129
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: EP01128519.4 PRIOR FILING DATE: 2001-11-29 NUMBER OF SEQ ID NOS: 312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: HELLENDOORN, Koen APPLICANT: BAKER, Matthew APPLICANT: WILLIAMS, Steven APPLICANT: CARR, Francis J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: CARR, Francis J.
TITLE OF INVENTION: T-CELL EPITOPES IN CARBOXYPEPTIDASE
                   ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 14
                                                              TYPE:
OTHER INFORMATION: Hair-binding peptide
                                                                             ENGTH: 12
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Pred. No. 7
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Pred. No. 6.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2e+02;
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and

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RESULT 15

US-11-074-473-48

US-11-074-473-48

Sequence 48, Application US/11074473

Publication No. US20050226839A1

APPLICANT: Huang, Xueying
APPLICANT: Wang, Hong

ITITLE OF INVENTION: Peptide-Based Conditioners and Colorants for Hair, Skin, and

ITITLE OF INVENTION: Nails

FILE REFERENCE: CL2296 US NA

CURRENT FILING DATE: 2005-03-08

PRIOR APPLICATION NUMBER: US/11/074,473

CURRENT FILING DATE: 2004-09-07

NUMBER OF SEQ ID NOS: 104

SOFTWARE: Patentin version 3.2

SEQ ID NO 48

LEUGTH: 12

TYPE: PRT

ORGANISM: Artificial Sequence

FEATURE:

FEATURE:

OTHER INFORMATION: Hair-binding peptide

US-11-074-473-48
Search completed: May 30, 2006, 10:24:41 Job time: 47 secs
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                                                                                                                                                            Query Match 35.1%; Score 26; DB 6; Length 12; Best Local Similarity 62.5%; Pred. No. 8.4e+02; Matches 5; Conservative 1; Mismatches 2; Indels
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1 TDMQAPTK 8
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                May 30, 2006, 10:22:25 ; Search time 5.66667 Seconds (without alignments) 29.481 Million cell updates/sec
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Match Length
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Gapop 10.0 , Gapext 0.5
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74
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Copyright (c) 1993 - 2006 Biocceleration Ltd
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US-10-516-079-64

US-10-516-079-94

US-11-118-524-5

US-10-538-066-625

US-10-538-066-15

US-10-538-066-15

US-11-176-182-52

US-10-473-6918-7

US-11-176-182-112

US-11-176-1877-2124

US-11-176-1877-2241

US-10-538-066-65

US-11-176-1877-1988

US-11-176-1877-1988

US-11-176-1877-1988
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 Sequence 64, Appl
Sequence 94, Appl
Sequence 5, Appli
Sequence 625, Appli
Sequence 4, Appli
Sequence 52, Appli
Sequence 152, Appli
Sequence 1987, Appli
Sequence 112, Appli
Sequence 112, Appli
Sequence 2124, Appli
Sequence 2240, Appli
Sequence 2241, Appli
Sequence 2241, Appli
Sequence 65, Appli
Sequence 66, Appli
Sequence 437, Appli
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|            | 45                | 44                | 43                | 42               | 41                | 40                | 9                 | 38                | 37               | 36                | 35                | J.                | 33                | 32                | 31                | 30                | 29                  | 28                 | 27                | 26                         |
|------------|-------------------|-------------------|-------------------|------------------|-------------------|-------------------|-------------------|-------------------|------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|---------------------|--------------------|-------------------|----------------------------|
|            | 17                | 17                | 17                | 17               | 17                | 17                | 17                | 17                | 17               | 17                | 17                | 18                | 18                | 18                | 18                | 18                | 18                  | 18                 | 18                | 18                         |
|            | 23.0              | -                 | -                 | 23.0             | _                 |                   |                   | 23.0              |                  |                   |                   |                   | 24.3              |                   | 24.3              |                   | 24.3                | •                  | •                 | 24.3                       |
|            | 14                | 14                | 13                | 13               | 11                | 10                | 10                | 9                 | 9                | ထ                 | 4                 | 15                | 15                | 12                | 11                | 11                | 10                  | 10                 | 10                | 10                         |
|            | 7                 | 7                 | 7                 | 7                | 7                 | σ                 | 9                 | 7                 | σ                | 7                 | 7                 | 7                 | 6                 | 7                 | σ                 | σ                 | 7                   | 7                  | g                 | σ                          |
| ALIGNMENTS | US-11-176-182-114 | US-11-176-182-63  | US-11-142-051-12  | US-11-298-718-44 | US-11-122-986-655 | US-10-538-066-240 | US-10-614-959-26  | US-11-122-986-263 | US-10-516-079-91 | US-11-122-986-30  | US-11-252-276-62  | US-11-122-986-725 | US-10-514-263-6   | US-11-122-986-570 | US-10-538-066-636 | US-10-538-066-479 | US-11-140-487A-1989 | US-11-140-487A-994 | US-10-538-066-574 | US-10-538- <b>0</b> 66-268 |
|            | Sequence 114, App | Sequence 63, Appl | Sequence 12, Appl | •                | ٠.                | •                 | Sequence 26, Appl | Sequence 263, App |                  | Sequence 30, Appl | Sequence 62, Appl | Sequence 725, App | Sequence 6, Appli | Sequence 570, App | Sequence 636, App | Sequence 479, App | Sequence 1989, Ap   | Sequence 994, App  | Sequence 574, App | -                          |

## RESULT 2 US-10-516-079-94 Sequence 94, Application US/10516079 Publication No. US20060088899A1 GENERAL INFORMATION: APPLICANT: ALVAREZ, Vernon L. APPLICANT: GONDA, Matthew A. TITLE OF INVENTION: Combination chemotherapy with chlorotoxin FILE REFERENCE: 51530-5006-WO CURRENT APPLICATION NUMBER: US/10/516,079 CURRENT FILING DATE: 2004-11-29 PRIOR APPLICATION NUMBER: US 60/406,033 APPLICANT: ALVAREZ, Vernon L. APPLICANT: GRIMES, Carol A. APPLICANT: GONDA, Matthew A. FILE OF INVENTION: Combination chemotherapy with chlorotoxin FILE REFERENCE: 51530-5006-WO CURRENT APPLICATION NUMBER: US/10/516,079 CURRENT FILING DATE: 2004-11-29 PRIOR APPLICATION NUMBER: US 60/406,033 PRIOR FILING DATE: 2002-08-27 PRIOR FILING DATE: 2002-08-27 PRIOR FILING DATE: 2002-05-31 PRIOR FILING DATE: 2002-05-31 NUMBER OF SEQ ID NOS: 95 COMPMEDE: Batalin version 3 2 RESULT 1 US-10-516-079-64 Sequence 64, Application US/10516079 Publication US20060088899A1 밁 Ś US-10-516-079-64 Query Match Best Local S Matches 5 SOFTWARE: PatentIn version 3.2 SEQ ID NO 64 LENGTH: 10 TYPE: PRT ORGANISM: Artificial sequence GENERAL INFORMATION: OTHER INFORMATION: Pep21-SCX8\_LEIQH sequence FEATURE: N 8 TDQQATTR 15 Similarity 5; Conserv TDQQMTKK 9 Conservative 35.1%; 62.5%; Score 26; DB Pred. No. 12; 1; Mismatches DB 12; ٠. ص Length 10; 0 Gaps

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APPLICANT: ALVAREZ, Vernon L.
APPLICANT: GRIMES, Carol A.
APPLICANT: GONDA, Matthew A.
APPLICANT: GONDA, Matthew A.
TITLE OF INVENTION: Combination chemotherapy with chl
FILE REFERENCE: 51530-5006-WO
CURRENT APPLICATION NUMBER: US/10/516,079
CURRENT FILING DATE: 2004-11-29
PRIOR APPLICATION NUMBER: US 60/406,033
PRIOR FILING DATE: 2002-08-27
PRIOR FILING DATE: 2002-05-31
PRIOR FILING DATE: 2002-05-31
NUMBER: US 60/384,171
NUMBER: OS 60/384,171
NUMBER: OS 60/384,171
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US-10-516-079-90
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; TYPE: PRT
; ORGANISM: homo sapiens
US-11-118-524-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: AFF-BP1 AS MEDIATOR OF PS3-DEPENDENT AND INDEPENDENT TUMOR TITLE OF INVENTION: SUPPRESSION AND USES THEREOF FILE REFERENCE: 19240-497US2 CURRENT APPLICATION NUMBER: US/11/118,524 CURRENT FILING DATE: 2005-04-29 PRIOR APPLICATION NUMBER: 60/610,506 PRIOR APPLICATION NUMBER: 60/610,506 PRIOR FILING DATE: 2004-09-15 NUMBER OF SEQ ID NOS: 22 SOFTWARE: Patentin version 3.3 SEQ ID NO 5
                                                                                                                                                                                                                                                                                                   Sequence 90, Application US/10516079 Publication No. US20060088899A1 GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
SOFTWARE: Pa
SEQ ID NO 90
LENGTH: 9
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Best Local :
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PRIOR APPLICATION NUMBER: US 60/384,171
PRIOR FILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 95
SOFTWARE: PatentIn version 3.2
SEQ ID NO 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: GU, WEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 9
TYPE: PRT
ORGANISM: Mesobuthus tamulus sindicus
FEATURE:
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Similarity 50.0%;
4; Conservative
                                        PatentIn version 3.2
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Pred. No. 1.1e+02;
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Pred. No. 5.2e+04;
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TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7,
TITLE OF INVENTION: Peptides and Compositions
FILE REFERENCE: 2060,015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 767
                                                                                                                        ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-538-066-4
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US-10-538-066-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-538-066-625
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US-10-538-066-625
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                                                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/10538066
Publication No. US20060094649A1
GENERAL INFORMATION:
APPLICANT: Epimmune Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 625, Application US/10538066 Publication No. US20060094649A1 GENERAL INFORMATION:
                                                             Matches
                                                                            Query Match
Best Local :
                                                                                                                                                                       SEQ ID NO 4
LENGTH: 11
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/432,017 PRIOR FILING DATE: 2002-12-10 NUMBER OF SEQ ID NOS: 767
                                                                                                                                                                                                                                                                  FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B
TITLE OF INVENTION: Peptides and Compositions
                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial sequence FEATURE:
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NIVI 8
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                             NINVL 5
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                                                             Conservative
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                                                                            27.0%; Score 20; 100.0%; Pred. No.
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Pred. No. 5.
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                                                             Mismatches
                                                                            DB 6;
2e+02;
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US-10-473-691B-7
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; TYPE: PRT
; ORGANISM: chimeric HIV-HRV
US-11-176-182-52
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US-11-176-182-52
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US-10-538-066-159
Sequence 7, Application US/10473691B
publication No. US20060099202A1
GENERAL INFORMATION:
APPLICANT: Nicolette, Charles A.
APPLICANT: Soltis, Daniel A.
APPLICANTION: IMMUNOSIOBULIN CONSTRUCT CONTAINING TUMOR SPECIFIC p53BP2 SEQUENC FILE REFERENCE: 02755/100K313-US1
CURRENT APPLICATION NUMBER: US/10/473,691B
CURRENT FILING DATE: 2003-09-29
PRIOR APPLICATION NUMBER: PCT/US02/10224
PRIOR APPLICATION NUMBER: PCT/US02/10224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Publication No. US20060094649A1
GENERAL INFORMATION:
APPLICANT: Epimmune Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7, and -B44 Tumor Associated Antigen
TITLE OF INVENTION: Peptides and Compositions
FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538.066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 767
SOFTWARE: Patentin version 3.2
SEQ ID NO 159
LENGTH: 9
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Publication No. US20060088549A1
GENERAL INFORMATION:
APPLICANT: ARNOLD, EDWARD
APPLICANT: FERSTANDIG ARNOLD, GAIL
TITLE OF INVENTION: CHIMERIC HIV VACCINE
FILE REFERENCE: PA04-701-S
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CURRENT FILING DATE: 2005-07-07
NUMBER OF SEQ ID NOS: 138
SOFTWARE: PatentIn version 3.3
SEQ ID NO 52
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 25.7%; Score 19; DB 6; Length 9;
Local Similarity 100.0%; Pred. No. 5.2e+04;
hes 4; Conservative 0; Mismatches 0; Indels
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Local Similarity 50.0%;
les 4; Conservative
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Pred. No. 2.6e+02;
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RESULT 11

US-11-176-182-112

Sequence 112, Application US/11176182

Sequence 112, DEPART OF SEQUENCE 
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TYPE: PRT
TYPE: PRT
CREAMISM: Artificial Sequence
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US-11-140-487A-1987
US-11-140-487A-1987
; Sequence 1987, Application US/11140487A
; Publication No. US20060093617A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: PRT; ORGANISM: hepatitis C virus US-11-140-487A-1987
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CURRENT FILING DATE: 2005-05-31
PRIOR APPLICATION NUMBER: EP 04012951.2
PRIOR FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: EP 04447239.7
PRIOR FILING DATE: 2004-10-28
PRIOR APPLICATION NUMBER: EP 05102441.2
PRIOR APPLICATION NUMBER: EP 05102441.2
PRIOR FILING DATE: 2005-03-25
PRIOR APPLICATION NUMBER: US 60/576,310
PRIOR APPLICATION NUMBER: US 60/576,310
PRIOR APPLICATION NUMBER: US 60/622,782
PRIOR FILING DATE: 2004-06-03
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Best Local
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 2278
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OTHER INFORMATION: P52 BP2
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Pred. No. 2.
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Pred. No. 5.
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RESULT 12
US-11-176-182-17
; Sequence 17, Application US/11176182
; Publication No. US20060088549A1
; GENERAL IMPORMATION:
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US-11-122-986-822
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FEATURE:
NAME/KEY: misc_feature
LOCATION: (12)...(12)
OTHER INFORMATION: Xaa can be any naturally occurring amino
US-11-176-182-112
        CURRENT FILING DATE: 2005-05-05
PRIOR APPLICATION NUMBER: 60/423,875
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,832
PRIOR APPLICATION NUMBER: 60/423,915
PRIOR APPLICATION NUMBER: 60/423,915
PRIOR FILING DATE: 2002-11-05
PRIOR PPLICATION NUMBER: 60/423,757
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,758
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,758
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/424,367
PRIOR APPLICATION NUMBER: 60/424,367
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/424,367
PRIOR FILING DATE: 2002-11-05
PRIOR FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,367
PRIOR FILING DATE: 2002-11-05
                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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LENGTH: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 822, A Publication No.
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Best Local :
                                                                                                                                                                                                                                                                         APPLICANT: DHARAMSI, AKIL
APPLICANT: VEDADI, WASOUD
TITLE OF INVENTION: ESSENTIAL NOVEL BACTERIAL POLYPEPTIDES
FILE REFERENCE: IPT-330.01
CURRENT APPLICATION NUMBER: US/11/122,986
CURRENT FILING DATE: 2005-05-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: ARNOLD, EDWARD
APPLICANT: FERSTANDIG ARNOLD, GAIL
TITLE OF INVENTION: CHIMERIC HIV VACCINE
FILE REFERENCE: PA04 - 701 - S
CURRENT FILLING DATE: 2005-07-07
NUMBER: OF SEQ ID NOS: 138
                                                                                                                                                                                                                                                                                                                                                             APPLICANT: EDWARDS, APPLICANT: DHARAMS APPLICANT: VEDADI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
APPLICATION NUMBER: 60/424,376
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Similarity 33.3%;
3; Conservation
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3; Conserv
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Pred. No. 3.8e+02;
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Pred. No. 3.5e+02;
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RESULT 15

US-11-140-487A-2125

¡ Sequence 2125, Application US/11140487A

¡ Publication No. US20060093617A1

¡ GENERAL INFORMATION:
¡ APPLICANT: Innogenetics N.V.
¡ TITLE OF INVENTION: Peptides for inducing a C
¡ FILE REFERENCE: 166
¡ CURRENT APPLICATION NUMBER: US/11/140,487A
; CURRENT FILING DATE: 2005-05-31
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US-11-140-487A-2124
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PRIOR FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,362
PRIOR FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/424,373
PRIOR FILING DATE: 2002-11-06
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 844
SOFTWARE: Patentin Ver. 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 2124, Application US/11140487A Publication No. US20060093617A1 GENERAL INFORMATION:
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Best Local Similarity 40.0%;
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SEQ ID NO 822
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PRIOR FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: EP 04447239.7
PRIOR FILING DATE: 2004-10-28
PRIOR APPLICATION NUMBER: EP 05102441.2
PRIOR FILING DATE: 2005-03-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 166
CURRENT APPLICATION NUMBER: US/11/140,487A
CURRENT FILING DATE: 2005-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 2004-10-29
PRIOR APPLICATION NUMBER: US 60/665,395
PRIOR FILING DATE: 2005-03-25
NUMBER OF SEQ ID NOS: 2278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/576,310 PRIOR FILING DATE: 2004-06-03 PRIOR APPLICATION NUMBER: US 60/622,782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Peptides for inducing a CTL and/or HTL response to Hepatitis C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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TYPE: PRT
ORGANISM: Enterococcus faecalis
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Pred. No. 4.5e+02;
2; Mismatches 4
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Pred. No. 3.8e+02;
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Peptides for inducing a

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Hepatitis

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PRIOR APPLICATION NUMBER: EP 04012951.2
PRIOR FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: EP 04447239.7
PRIOR APPLICATION NUMBER: EP 04447239.7
PRIOR PRILING DATE: 2004-10-28
PRIOR PILING DATE: 2004-10-28
PRIOR FILING DATE: 2004-06-03
PRIOR FILING DATE: 2004-06-03
PRIOR FILING DATE: 2004-06-03
PRIOR FILING DATE: 2004-06-03
PRIOR APPLICATION NUMBER: US 60/665,395
PRIOR FILING DATE: 2004-06-03
PRIOR FILING DATE: 2005-03-25
PRIOR FILING DATE: 2005-03-25
PRIOR APPLICATION NUMBER: US 60/665,395
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Maximum Match 100%
Listing first 45 s
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Gapop 10.0 , Gapext 0.5
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l: /EMC_Celerra_SID:
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/EMC_Celerra_SIDS3/ptodata/2/iaa/6_COMB.pep:*
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/EMC_Celerra_SIDS3/ptodata/2/iaa/PCTUS_COMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*
/EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*
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  GenCore version 5.1.8
(c) 1993 - 2006 Biocceleration Ltd
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US-09-815-567B-2
US-09-813-567B-2
US-09-016-061-88
US-09-016-061-88
9523254-6
US-09-641-528B-48206
US-09-878-281A-244
US-09-878-281A-244
US-09-878-281A-244
US-09-681-1366
US-10-000-986A-144
US-09-641-528B-4922
US-09-641-528B-4922
US-09-641-528B-49305
US-09-641-528B-49305
US-09-641-528B-493032
US-09-641-528B-49303
US-09-641-528B-493032
US-09-641-528B-493032
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US-09-641-528B-4923
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Sequence 88, Appli
Sequence 88, Appli
Sequence 88, Appl
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Sequence 166, App
Sequence 166, App
Sequence 166, App
Sequence 166, App
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56, Appl
492, Ap
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45697, A
49305, A
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50384, A
50384, A
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50384, A
504, Appl
505, Appl
504, Appl
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| 4 4 5<br>5 4 6                               | 4 4 4<br>2 2 1   | 39                                    | 37<br>38                                     | 35<br>6                                      | 3<br>4<br>4                                  | 31<br>32                                    | 30<br>30                                    | 27<br>28             |
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| 206  | 2 2 2 6 6  | 26                                    | 26<br>26                                     | 26<br>26                                     | 26<br>26                                     | 26<br>26                                    | 26<br>6                                     | 26                   |
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| US-09-641-528B-47147<br>US-09-641-528B-47208 | US-08-837-941-28<br>US-09-148-712-23<br>US-09-535-364-10 | US-09-620-091-172<br>US-08-321-668-28 | US-09-641-528B-42043<br>US-09-641-528B-45698 | US-09-641-528B-26893<br>US-09-641-528B-34845 | US-09-641-528B-13785<br>US-09-641-528B-26703 | US-09-641-528B-4856<br>US-09-641-528B-13626 | US-09-641-528B-26913<br>US-09-641-528B-4810 | -09-641              |
| Sequence<br>Sequence                         | Sequence<br>Sequence                                     | Sequence<br>Sequence                  | Sequence<br>Sequence                         | Sequence<br>Sequence                         | Sequence<br>Sequence                         | Sequence<br>Sequence                        | Sequence                                    | Sequence<br>Sequence |
| 47147, A<br>47208, A                         | 28, Appl<br>23, Appl<br>10, Appl                         | 172, App<br>28, Appl                  | 42043, A<br>45698, A                         | 26893, A<br>34845, A                         | 13785, A<br>26703, A                         | 4856, Ap<br>13626, A                        | 26913, A<br>4810, Ap                        | 13808, A<br>26799, A |

# ALIGNMENTS

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CURRENT APPLICATION NUMBER: US/09/526,193A
CURRENT FILING DATE: 2000-03-15
PRIOR APPLICATION NUMBER: 60/124,702
PRIOR FILING DATE: 1999-03-15
PRIOR PRILING DATE: 1999-03-15
PRIOR APPLICATION NUMBER: 60/138,048
PRIOR FILING DATE: 1999-06-17
PRIOR APPLICATION NUMBER: 60/139,600
PRIOR FILING DATE: 1999-06-17
PRIOR APPLICATION NUMBER: 60/151,977
PRIOR FILING DATE: 1999-09-01
NUMBER OF SEO ID NOS: 287
SOFTWARE: FRSTSEO for Windows Version 4.0
SEQ ID NO 65
LENGTH: 10
; Sequence 2, Application US/08851567B
; Patent No. 6528484
; GENERAL INFORMATION:
   APPLICANT: Ensign, Jerald C
   APPLICANT: Bowen, David J
   APPLICANT: Petell, James
   APPLICANT: Fatig, Raymond
   APPLICANT: Schoonover, Sue
   APPLICANT: ffrench-Constant, Richar
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US-09-526-193A-65
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Best Local (
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APPLICANT: Brooks-Wilson, Angela R.
APPLICANT: Pimstone, Simon N.
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TITLE OF INVENTION: METHODS AND REAGENTS FOR MODULATING
TITLE OF INVENTION: CHOLESTEROL LEVELS
FILE REFERENCE: 50110/002005
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                    l Similarity
6; Conserv
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Ensign, Jerald C
Bowen, David J
Petell, James
Patig, Raymond
Schoonover, Sue
ffrench-Constant, Richard
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Pred. No. 64;
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Gaps

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APPLICANT:

Rocheleau, Thomas A. Blackburn, Michael B

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RESULT 3
US-09-309-922A-88
; Sequence 88, Application US/09339922A
; Patent No. 6531580
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                                                                                                                                                                                                                                  Query Match
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APPLICATION NUMBER: US 08/063,615
FILING DATE: 18-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/395,497
FILING DATE: 28-FEB-1995
PRIOR APPLICATION NUMBER: US 60/007,255
PRICING DATE: 06-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 608-251-9166
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: protein FRAGMENT TYPE: N-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 0 FILING DATE: 28-FEB-1996 PRIOR APPLICATION DATA:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/851,567B
FILING DATE: 05-MAY-1997
CLASSIFICATION: 514
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
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APPLICANT: Ciche, Todd A.
APPLICANT: Sukhapinda, Kitisri
TITLE OF INVENTION: Insecticidal Protein Toxins From Photorhabdus
NUMBER OF SEQUENCES: 88
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dow AgroSciences Patent Department
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 08/705,484 FILING DATE: 28-AUG-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 608-251-5000
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                                                                                                                                                                                  33.5%;
Local Similarity 53.8%;
les 7; Conservaring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 9330 Zionsville Road CITY: Indianapolis
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Strickland, James A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nicholas :
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Query Match
Best Local Similarity
Thes 4; Conserve
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LENGTH: 9
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Query Match
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APPLICANT: Wu, Herren
TITLE OF INVENTION: Anti-AlphaV Beta3 Recombinant Human Antibodies, Nucleic
TITLE OF INVENTION: Acids Encoding Same and Methods of Use
FILE REFERENCE: P-1X 3536
CURRENT APPLICATION NUMBER: US/09/339,922A
CURRENT FILING DATE: 1999-06-24
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                     FILING DATE: 30-JAN-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/791,391
FILING DATE: 30-JAN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-IX 2965
                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9901
TELEPAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 88:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: complementarity determining region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PA PC COMPATIBLE
COMPUTER: PATENT PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Huse, William D.

APPLICANT: Glaser, Scott M.

TITLE OF INVENTION: Anti-Alpha V Beta 3 Recombinant Human

TITLE OF INVENTION: Antibodies, Nucleic Acids Encoding Same and Methods of Use
                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 4370 La Jo
CITY: San Diego
STATE: California
                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/016,061
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                                                                              amino acid
GY: linear
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4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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Pred. No. 5e+05;
    Score 28;
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APPLICANT: Sette, Alessandro
APPLICANT: Southwood, Scott
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
APPLICANT: Chesnut, Robert
APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 48206
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RESULT 7
US-08-836-075A-166
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                                                                                                                                                                                                                 JS-09-641-528B-48206
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Matches 4; Conserval
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GENERAL INFORMATION:
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                                                                                                                                          Matches
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                                                                                                                                                                                                                        FEATURE:
OTHER INFORMATION: Peptide Derived from Human Papillomavirus
                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                    ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/07/247,017 FILING DATE: 20-SEP-1988 PRIOR APPLICATION DATA:
                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 102,180 FILING DATE: 29-SEP-1987
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Pred. No. 1.9e+02;
3; Mismatches 3
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2; Mismatches
                                                                                                                                                           Score 28; DB 3;
Pred. No. 2.2e+02;
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                                                   GENERAL INFORMATION:
APPLICANT: Innogenetics N.V.
TITLE OF INVENTION: New sequences of hepatitis C virus genotypes for diagnosis, prophy
TITLE OF INVENTION: and therapy
FILE REFERENCE: 35
CURRENT APPLICATION NUMBER: US/09/878,281A
CURRENT FILING DATE: 2001-06-12
NUMBER OF SEQ ID NOS: 284
SOFTWARE: PatentIn version 3.1
SEQ ID NO 244
LENGTH: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 166, Application US/08836075A Patent No. 6180768
                                                                                                                                                                                                                                                                      Sequence 244, Application US/09878281A Patent No. 6762024
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Best Local Similarity
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ZIP: 77210-4433
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC_DOS/MS-DOS
SOFTWARE: Microsoft Word 6.0 / AS(
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: PCT/EP95/04155
PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: EP 94870166.9
PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: EP 95870076.7
PRILING DATE: 28 Jun 1995
ATTORNEY/AGENT INFORMATION: NAME: KAMMERER, PATRICIA A.
REGISTRATION NUMBER: 29,775
09-878-281A-244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: INFORMATION FOR SEQ ID NO: 1 SEQUENCE CHARACTERISTICS:
            LENGTH: 13
TYPE: PRT
ORGANISM: hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: 1i
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: MAERTENS, APPLICANT: STUYVER,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: HOUSTON STATE: TEXAS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
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                                                                                                                                                                                                                                                                                                                                                                                        VRTGNQSRCW 10
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50.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 27; DB 2;
Pred. No. 2.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 13;
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Best Local Similarity
"hes 5; Conserv?
                                         US-10-000-986A-144
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US-10-000-986A-144
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TYPE: PRT
CORGANISM: hepatitis C virus
US-09-851-138C-166
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                                                                               SOFTWARE: JPE
SEQ ID NO 144
LENGTH: 14
TYPE: PRT
Query Match
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APPLICANT: INNOGENETICS N.V.
APPLICANT: INNOGENETICS N.V.
TITLE OF INVENTION: New sequences of hepatitis C virus genotypes and their use as
TITLE OF INVENTION: prophylactic, therapeutic and diagnostic agents.
FILE REFERENCE: 2551-106
CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: EP 94870166.9
PRIOR FILING DATE: 1994-10-21
PRIOR FILING DATE: 1994-10-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 166, Appl
Patent No. 6974864
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                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/302,277
PRIOR PILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: US 60/298,698
PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: US 60/293,574
PRIOR FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Benjanin, Stephane
APPLICANT: Tanaka, Hiroaki
TITLE OF INVENTION: HUMAN CDNAS AND PROTEINS AND USES THEREOF
FILE REFERENCE: G-091US09DIV
                                                                                                                                                                    NUMBER OF SEQ ID NOS: 228
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SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: EP 95870076.7 PRIOR FILING DATE: 1995-06-28
                                                      ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31.8%;
Local Similarity 50.0%;
es 5; Conservation
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                                                                                                                                              JPatent
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31.8%;
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Pred. No. 2.9e.
1; Mismatches
Score 27;
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Pred. No. 2.9e+02;
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DB 3;
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2.9e+02;
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Length 14;
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                                                                                                                                                                                                                               PCT-US95-04018-56
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                                                                                                                                                                                             Sequence 56, Application PC/TUS9504018 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: DELUCA, MARK
REGISTRATION NUMBER: 3,729
REFERENCE/DOCKET NUMBER: CC
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                   TITLE OF INVENTION: Tumor Necrosis Factor Inhibitors
                                                                                                                                    APPLICANT:
                                                                                                                                                                 APPLICANT: Heavner, George A. APPLICANT: Kruszynski, Marian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Heavner, TITLE OF INVENTION:
                                                                                       CORRESPONDENCE ADDRESS:
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                          STREET: One Liberty Place CITY: Philadelphia
                                                     ADDRESSEE: Woodcoo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: One Liberty Place CITY: Philadelphia
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                  Local
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6; Conservative
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              Pennsylvania
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: USA
                                                                                                                                Kruszynski, Marian
Mervic, Miljenko
Weber, Robert W.
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(215) 56
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e Liberty Place 46th Floor
                                                                        Woodcock Washburn Kurtz Mackiewicz &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     568-3439
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Tumor Necrosis Factor Inhibitors
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Pred. No. 3
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                                            46th Floor
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3.4e+02;
7;
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MSSOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:

PC-DOS/MS-DOS

PCT/US95/04018

CLASSIFICATION: APPLICATION NUMBER: ZIP: 19403 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk

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APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
SEQ ID NOS: 51505
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4922
LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; MOLECULE TYPE: peptide
PCT-US95-04018-56
                                                           US-09-641-528B-4922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4922, Application US/09641528B Patent No. 7026443
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Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
SEQUENCE TRACTER 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/221,580
FILING DATE: 01-APR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/221,583
FILING DATE: 01-APR-1994
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/221,581
APPLICATION UMBER: US 08/221,581
APPLICATION UMBER: US 08/221,581
FILING DATE: 01-APR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                            OTHER INFORMATION: Peptide Derived from Human Papillomavirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (215) 568-3100
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Local Similarity 46.2%;
hes 6; Conservaring
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: DeLuca, Mark
REGISTRATION NUMBER: 33,229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HLPQPVSTRSQHT 13
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  30.6%;
55.6%;
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Pred. No. 3.4:
0; Mismatches
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  Score 26;
Pred. No.
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Patent No. 7026443
GENERAL INFORMATION:
APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
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                                      US-09-641-528B-26852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 15
US-09-641-528B-26852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Peptide Derived from Human Papillomavirus US-09-641-528B-13784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-641-528B-13784
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 13784, Application US/09641528B Patent No. 7026443 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 13784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                              SEQ ID NO 26852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 26852, Application US/09641528B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Query Match
                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/641,528B CURRENT FILING DATE: 2000-08-15 PRIOR APPLICATION NUMBER: US 60/172,705 PRIOR FILING DATE: 1999-12-10 NUMBER OF SEQ ID NOS: 51505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/641,528B CURRENT FILING DATE: 2000-08-15 PRIOR APPLICATION NUMBER: US 60/172,705 PRIOR FILING DATE: 1999-12-10
                                                                                                                                                                                                                                                                    TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS FILE REFERENCE: 2060.0100001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS FILE REFERENCE: 2060.0100001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0
                                                   FEATURE: OTHER INFORMATION: Peptide Derived from Human Papillomavirus
                                                                                                            LENGTH: 9
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence FEATURE:
                                                                                       ORGANISM: Artificial Sequence
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Southwood, Scott
Chesnut, Robert
Celis, Esteban
Grey, Howard
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Grey, Howard
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Chesnut, Robert
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Pred. No. 5e+05;
   Score 26;
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|  | Search completed: May 30, 2006, 11:11:06  Job time : 23.3333 secs | Best Local Similarity 57.1%; Pred. No. 5e+05; Matches 4; Conservative 2; Mismatches 1; Indels 0; Gaps 0;  Qy 9 EQQATTW 15       : : |
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| 45                 | 44                 | 43                 | 42                | 41                 | 40                 | 39                 | 38                 | 37                | 36                 | 35     | υ<br>4             | υ<br>33            | 32                 | 31                 | 30                 |
|--------------------|--------------------|--------------------|-------------------|--------------------|--------------------|--------------------|--------------------|-------------------|--------------------|--------|--------------------|--------------------|--------------------|--------------------|--------------------|
| 17                 | 17                 | 17                 | 17                | 17                 | 17                 | 17                 | 17                 | 17                | 17                 | 17     | 17                 | 17                 | 18                 | 18                 | 18                 |
| 20.0               | 20.0               | 20.0               | 20.0              | 20.0               | 20.0               | 20.0               | 20.0               | 20.0              | 20.0               | 20.0   | 20.0               | 20.0               | 21.2               | 21.2               | 21.2               |
| 11                 | 11                 | 11                 | 11                | 10                 | 10                 | 10                 | 9                  | œ                 | œ                  | 7      | σ                  | տ                  | 15                 | 15                 | 15                 |
| N                  | N                  | N                  | N                 | N                  | N                  | N                  | N                  | N                 | N                  | N      | N                  | N                  | N                  | N                  | N                  |
| A61512             | C49037             | B49037             | H54346            | S39030             | A60722             | S66248             | PT0247             | 157018            | PT0030             | S33246 | A31263             | PT0580             | A53594             | A49177             | PA0058             |
| variant surface gl | TcR gamma V-J regi | TcR gamma V-J regi | pyruvate synthase | lysyl-bradykinin - | cryptic fimbrial p | processing enzyme, | Ig heavy chain CRD | gene Cftr protein | inulinase (EC 3.2. | ~      | dihydrofolate redu | T-cell receptor be | calnexin - mouse ( | 22K protein pl, mi | protein QF200022 - |

# ALIGNMENTS

RESULT 1

R;Frebort, I.; Tamaki, H.; Ishida, H.; Pec, P.; Luhova, L.; Tsuno, H.; Halata, M.; Asano Bur. J. Biochem. 237, 255-265, 1996
A;Title: Two distinct quinoprotein amine oxidases are induced by n-butylamine in the myc A;Reference number: S71303; MUID:96203933; PMID:8620882
A;Accession: S71304

A;Cross-references: UNIPROT:Q7M504; UNIPARC:UPI000017B3B7 C;Keywords: copper binding; monomer; oxidoreductase; quinoprotein; topaquinone

A;Molecule type: protein A;Residues: 1-11 <FRE>

amine oxidase (copper-containing) (EC 1.4.3.6) II - Aspergillus niger (fragment) C;Species: Aspergillus niger (C;Species: Aspergillus niger (C;Date: 12-Feb-1998 #sequence\_revision 01-May-1998 #text\_change 09-Jul-2004 C;Accession: S71304 C;Accession: S71304 H; Ishida, H; Pec, P; Luhova, L; Tsuno, H; Halata, M; R;Frebort, I; Tamaki, H; Ishida, H; Pec, P; Luhova, L;

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RESULT 3
I41235
                                                                                                                                                                                                                                                                                                C;Accession: PA0062
R;Chow, L.P.; Fukaya, N.; Sugiura, Y.; Ueno, Y.; Tabuchi, K.; Tsugita, A.
submitted to JIPID, October 1994
A;Description: Two dimensional polyacrylamide gel electropheresis of Fusarium sporotrich
A;Reference number: PA0051
A;Accession: PA0062
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 2
PA0062
                                                                                                                                                                                                                                                                                                                                                                                                                        fumarate hydratase (BC 4.2.1.2) - fungus (Fusarium sporotrichioides) (fragment)
C;Species: Fusarium sporotrichioides
C;Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 09-Jul-2004
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Best Local Similarity
                                                                                                                                               y Match
25.9%; Score 22; DB 2; Length 15;
Local Similarity 55.6%; Pred. No. 1.4e+03;
hes 5; Conservative 0: Miamoral-
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                                                                                                                 4 SPVRTEQQA 12
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                                                                            SQTRTESDA 9
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glucocerebrosidase - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 02-Jul-1996 #sequence_revisio:
C;Accession: I52980; I65971
                                                                                                                                                                                                                                                                                                                                 C;Species: Homo sapiens (man)
C;Date: 02-Jul-1996 #text_change 05-Nov-1999
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 05-Nov-1999
C;Accession: 152980; 165971
R;Reiner; O.; Wigderson, M.; Horowitz, M.
DNA 7, 107-116, 1988
DNA 7, 107-116, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        glutamine-tRNA ligase (EC 6.1.1.18) - Escherichia coli (fragment) C;Species: Bscherichia coli (c;Species: Ascherichia coli (c;Date: 31-May-1996 #sequence_revision 31-May-1996 #text_change (c;Date: 31-May-1996 #sequence_revision 31-May-1996 #text_change (c;Accession: I41235 R;Hoben, P.; Uemura, H.; Yamao, F.; Cheung, A.; Swanson, R.; Sumr Fed. Proc. 43, 2972-2976, 1984
                                                                                                                                                                                                          A;Status: preliminary; translated from GB/EMBL/DDBJA;Molecule type: DNAA;Residues: 1-11 <RES>
                                                                                                                                                                                                                                                                       A;Title: Structural analysis of the human glucocerebrosidase A;Reference number: I52980; MUID:88195776; PMID:3359914 A;Accession: I52980
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J. Exp. Med. 173, 19-24, 1991
A;Title: A myelin basic protein peptide is
A;Reference number: PHO135; MUID:91086843;
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C;Accession: PH0138
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                                                                              A;Cross-references: UNIPARC:UPI000016A981; GB:M18917; NID:g183025; PIDN:AAA35879.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Accession: PH0138
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-12 < RES>
                                                                                                        ;Molecule type: DNA
;Residues: 1-11 <RE
                                                                                                                                           Status: preliminary; translated
                                                                                                                                                                   Accession: I65971
                                                                                                                                                                                      ;Cross-references: UNIPARC:UPI000016A981; GB:M18916; NID:g183023; PIDN:AAA35878.1;
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Keywords: aminoacyl-tRNA synthetase; ligase; protein biosynthesis
                       Local
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  Similarity
4; Conserv
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3; Conservative
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75.0%;
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                     100.0%;
                                         23.5%;
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Pred. No. 1.8e
1; Mismatches
                     Score 20;
Pred. No

    Mismatches

                                                                                                                                               from GB/EMBL/DDBJ
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Pred. No. 1.7e+03
  Mismatches
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1.8e+03;
0;
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  2.3e+03;
hes 0;
                                         DB 2;
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                                         Length 11;
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C;Species: 1
                                                                              R;Anastasi, A.; Bertaccini, G.; Cei, J.M.; De Caro, G.; Erspamer, V.; Impicciatore, Br. J. Pharmacol. 37, 198-206, 1969
Br. J. Pharmacol. 37, 198-206, 1969
A;Title: Structure and pharmacological actions of phyllocaerulein, a caerulein-like A;Reference number: A61357; MUID:70005484; PMID:5824931
A;Accession: A61357
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A61357
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                     A; Molecule type: protein A; Residues: 1-9 < ANA>
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                                                            A;Status: preliminary
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phyllocaerulein - Sauvage's leaf frog
C;Species: Phyllomedusa sauvagei (Sauvage's leaf frog)
C;Date: 09-Sep-1994 #sequence_revision 09-Sep-1994 #text_change 09-Jul-2004
C;Accession: A61357
                                                                                                                                                                                                                                                                                                                                                                                                     R;Fiske, M.J.; Tobey-Fincher, K.L.; Fuchs, R.L.
J. Gen. Microbiol. 136, 2377-2383, 1990
A;Title: Cloning of two genes from Bacillus circulans WL-12 which encode 1,3-beta-glucans A;Reference number: A60763; MUID:91178514; PMID:2127800
A;Accession: B60763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Thermus sp.
C;Date: 18-Feb-1994 #sequence_revision 19-Apr-1996 #text_change 07-May-1999
C;Accession: S39012
                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT:Q7M110;
C;Comment: This bacillus produces up
C;Keywords: glycosidase; hydrolase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   endo-1,3-beta-glucanase (EC 3.2.1.-), 40k - Bacillus circulans (strain WL-12) C;Species: Bacillus circulans C;Species: 14-May-1993 #sequence_revision 14-May-1993 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Freeman, S.A.; Peek, K.; Prescott, M.; Daniel, R. Biochem. J. 295, 463, 469, 193
A;Tille: Characterization of a chelator-resistant proteinase A;Reference number: S39012; MUID:94058984; PMID:8240244
                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: protein A; Residues: 1-15 < FIS>
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A;Molecule type: protein
A;Residues: 1-15 <FRE>
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Note: 13-Ala was also
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Pred. No. 3.2e
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Pred. No. 3
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p to six different 1,3-beta-glucanases
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UNIPROT: Q7LZC4;

UNIPARC: UPI000017668D

caerulein-like

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RESULT 11
A49037
TcR gamma
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R;Mehansho, H.; Ann, D.K.; Butler, L.G.; Rogler, J.; Carlson, D.M.
J. Biol. Chem. 262, 12344-12350, 1987
J. Biol. Chem. 262, 12344-12350, 1987
A;Title: Induction of proline-rich proteins in hamster salivary glands by isoproterenol A;Reference number: A92611; MUID:87308247; PMID:3040740
A;Accession: A29806
                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 10
A29806
cidic proline-rich protein HP43b - golden hamster (fragment)
acidic proline-rich protein HP43b - golden hamster (fragment)
C;Species: Mesocricetus auratus (golden hamster)
C;Date: 19-May-1989 #sequence_revision 19-May-1989 #text_chan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Superfamily: gastrin
C;Keywords: amidated carboxyl end; antihypertensive; neuropeptide; pyroglutamic acid;
F;1/Modified site: pyrrolidone carboxylic acid (cln) #status experimental
F;4/Binding site: sulfate (Tyr) (covalent) #status experimental
F;10/Modified site: amidated carboxyl end (Phe) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: protein
A;Residues: 1-10 <ANA>
A;Residues: 1-10 <ANA>
A;Cossidues: 1-10 <ANA>
A;Cossidues: 100 <ANA>
A;Cossidues: 100 <ANA>
A;Comment: The last five amino acids and the carboxyl terminal amide group of t
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Arch. Biochem. Biophys. 125, 57-68, 1968
A;Title: Isolation and amino acid sequence of caerulein,
A;Reference number: A61337; MUID:68238534; PMID:5649531
A;Accession: A61337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Superfamily: gastrin C;Keywords: amidated carboxyl end; neuropeptide; pyroglutamic acid; skin; F;1/Mcdified site: pyrrolidone carboxylic acid (Gln) #status experimental F;3/Binding site: sulfate (Tyr) (covalent) #status experimental
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A61337
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C;Date: 05-Aug-1994 #sequence_revision 05-Aug-1994 #text_change 07-May-1999
C;Accession: A61337
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Pred. No. 3.4e+03;
1; Mismatches 1
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Pred. No. 3.1e+03
1; Mismatches
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R;Sterjiades, R.; Pelmont, J.
Appl. Environ. Microbiol. 55, 340-347, 1989
A;Title: Occurrence of two different forms of protocatechuate
A;Reference number: A60458; MUID:89245845; PMID:2541659
A;Accession: A60458
                                                                                                                                                                                                                                                                                                protocatechuate 3,4-dioxygenase (EC 1.13.11.3) alpha chain - Moraxella sp. (strain N;Alternate names: protocatechuate oxygenase C;Species: Moraxella sp. C;Species: Moraxella sp. C;Date: 20-Feb-1993 #sequence_revision 20-Fab-1993 #text_change 07-May-1999 C;Accession: A60458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   43.2K bile stone protein - unidentified bacterium (fragment) C;Species: unidentified bacterium C;Date: 07-Feb-1997 #sequence_revision 07-Feb-1997 #text_cha C;Accession: E58502 R;Binette, J.P.; Binette, M.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Mus musculus (house mouse)
C;Date: 21-Jan-1994 #sequence_revision 18-Nov-1994 #text_change 05-Nov-1999
C;Accession: A49037
R;Ezquerra, A.; Wilde, D.B.; McConnell, T.J.; Sturmhofel, K.; Valas, R.B.;
                                                                             A;Molecule type: protein
A;Residues: 1-13 <STE>
A;Cross-references: UNIPARC:UPI000017A9FE
A;Note: two forms P and G of the alpha sul
C;Keywords: iron; oxidoreductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:Q7MIDO; UNIPARC:UPI000017A8D4
A;Experimental source: human bile with stones
A;Note: a secondary sequence DVKIGVAGS was also found
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Description: The proteins A;Reference number: A58501 A;Accession: E58502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eur. J. Immunol. 22, 491-498, 1992
A;Title: Mouse autoreactive gamma/delta T cells. II. Molecular characterization of the
A;Reference number: A49037; MUID:92164730; PMID:1311262
A;Accession: A49037
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A;Note: sequence extracted from NCBI backbone (NCBIN:90637, NCBIP:90641)
                                                                                                                                                                                                                                                                                                                                                                                                                  A60458
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A;Molecule type: DNA
                                      Query Match
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Pred. No. 4.1e+03;
2; Mismatches 3;
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Pred. No. 3.8e+03
1; Mismatches
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Pred. No. 3.4e+03;
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gallbladder st
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Superioxide dismutase (EC 1.15.1.1) 1 (NI) - Streptomyces coelicolor (Fragment)
C.Species: Streptomyces coelicolor
C.Date: 11-Mar 1998 #sequence_revision 17-Apr-1998 #text_change 07-May-1999
C.Accession: S74128
R.Kim, B.J.; Kim, H.P.; Hah, Y.C.; Roe, J.H.
Eur. J. Biochem. 241, 178-185, 1996
A.Fitle: Differential expression of superoxide dismutases containing Ni and Fe/Zn in Str A.Reference number: S74128; MUID:97054607; PMID:8898904
A.Rocession: S74128
A.Rolecule type: protein
A.Roses-references: UNIPARC:UPI000017AE10
A.Experimental source: ATCC 10147
C.Function:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ribosomal protein S14 - Mycobacterium bovis (fragments)
C;Species: Mycobacterium bovis
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 13-Jan-1995
C;Accession: S36887
R;Ohara, N.; Kimura, M.; Higashi, Y.; Yamada, T.
FEBS Lett. 331, 9-14, 1993
A;Title: Isolation and amino acid sequence of the 30S ribosomal protein S19 from Mycobac A;Reference number: S36887; MUID:94009653; PMID:8405418
A;Accession: S36887
A;Status: preliminary
Search completed: May 30, Job time : 13 secs
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                                                                                                                                                                                                                                                              A; Description: catalyzes the dismutation of 2 molecules of peroxide radical to dioxygen C; Keywords: metalloprotein; nickel; oxidoreductase; tetramer
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A;Residues: 1-2;3-12;13 <OHA>
A;Cross-references: UNIPARC:UPI000061DAD; UNIPARC:UPI000011EC65; UNIPARC:UPI000017AD47
C;Keywords: protein biosynthesis; ribosome
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Best Local Similarity 37.5%;
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Local Similarity 60.0%;
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                  2006, 10:13:20
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Pred. No. 4.5e+03;
0; Mismatches 2
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Pred. No. 4.1e+03;
4; Mismatches 1
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Maximum Match
Listing first
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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y length: 15
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85
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1: uniprot_sprot:*
2: uniprot_trembl:*
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Q71GS6 9HYME
Q71GS6 AZOVI
Q65CH4 9GEMI
RIPL LUFCY
Q9T2WO YEAST
O71066 9PARA
Q86D10 TRYCR
Q8TDAB HUMAN
Q6JDG0 CANFA
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Q6LBJO MOUSE
Q7M504 ASPNG
Q47602 ECOLI
Q50117 MYCLE
Q7M423 FUSSP
Q9T339 9PRIM
CWP06 PHAVU
Q798K5 STRLI
Q71GM6_9HYME
Q6LDN2_BACST
Q71GV6_9HYME
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Q7M110_BACCI
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Q5ed33
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Q71G86
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  Q9t2w0
Q71036
Q86d30
Q8tda8
Q6jdg0
Q9eqv3
Q718t2
Q718t2
Q71gm6
Q61dn2
Q71fm6
Q61dn2
Q71h38
Q71h38
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Q47602 escherichia
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Q6lbj0 mus musculu
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oryza sativ
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fusarium sp
                                                                                                                tomato leaf
Q6LBJO_MOUSE
ID Q6LBJO, MOUSE PRELIMINARY; PRT; 11 AA.
AC Q6LBJO;
DT 05-JUL-2004, integrated into UniProtKB/TrEMBL.
DT 07-FEB-2006, entry version 1.
DT 07-FEB-2006, entry version 8.
DE Topoisomerase I (Fragment).
GN Name=Top1; Synonyms=TOP;
OS Mus musculus (Mouse).
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| RESULT 2 |           |            | Query Match<br>Best Local<br>Matches                          | SEQUENCE              | EMBL;           |  | populations.";<br>J. Bacteriol. |                        | Datta S., Shirai           |  |             |      |            |                               |                         |            |        | ~~                  | F9          |            | 4 4 4<br>W 4 7   | 41<br>42                     | 0 6 6<br>0 6              | ນ ເມ<br>ເປີ 0<br>7 0 | , ω <sub>Φ</sub> | ) W W                        |  |
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|          |           |            | 28.<br>57.<br>tive  | 273                   | 8486            | O D                                      | iĭ                              | nn S., Be<br>genotypes | Naka                       | vadny  | .11.3210-32 |      |            | :teria; Epsil<br>Helicobacter | (Cam                    | (Fragment) | 5 <    | ted:                | PRELIMINARY |            | 12   | 10                           | 100                       | 500                  |                  | 15                           |  |
|          |           |            | 120   | MW;                   | 1;              | eat i                                    | 3218                            | of of                  | Zav                        | val.   | -0-3        |      |            | )<br>Eg                       | )<br>1<br>V             | nent       | is:    | into                | ξ;          |            | 222  |                              |                           |                      |                  |                              |  |
|          |           |            | Score 24; DB 2; Len<br>Pred. No. 4.1e+03;<br>1; Mismatches 2; | CAD126337B133DC6 CRC6 | -; Genomic_DNA. | Commons Attributi                        | 3218 (2000) .                   | 710                    | lly R., Segal              | k., verapatino 8., s<br>lez Y., Mistry R.S., ( | 18.2000;    | 1    |            | onproteobacteria;             | (Campylobacter pylori). | t).        | n 1.   | o UniProtKB/TrEMBL. | PRT; 10 AA. | ALIGNMENTS | Q53EA1 9NEOB<br>Q7M1D0_9BACT<br>Q98Z40_9HIV1                   | Q9QVE5_9MURI<br>P84487_LITCE | COCIN PHACN<br>SC41_TITCA | CAELZ_LITSP          | Q71H00_9HYME     | Q9R531 9DEIN<br>Q8JE81 9HIV1 |  |
|          |           |            | Length 10; ; Indels 0; Gaps 0                                 | 64;                   |                 | /www.uniprot.org/t<br>n-NoDerivs License |                                 | m different hum        | Wong B.C., O. Schneider R. | ilman R.H., Yu                                 | :           |      |            | Campylobacterales;            |                         |            |        |                     |             |            | Q5Jear ereucheroda<br>Q7m1d0 unidentifie<br>Q98z40 human immun |                              |                           | P62541 litoria ere   | andrena          | ther                         |  |
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RESULT
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PROTEIN SEQUENCE.

MEDLINE=96203933; PubMed=8620882;

Frebort I., Tamaki H., Ishida H., Pec P., Luh
Frebort I., Asano Y., Kato Y., Matsushita K.,
                                                                                           NON TER
                                                                                                                                                                                                                                                                                                  "Two distinct quinoprotein amine oxidases are induced by n-butylamine in the mycelia of Aspergillus niger AKU 3302. Purification, characterization, CDNA cloning and sequencing."; Eur. J. Blochem. 237:255-265(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Fungi; Ascomycota; Eurotiales; Trichocomaceae; mi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-DEC-2003, integrated into UniProtKB/TrEMBI.
15-DEC-2003, sequence version 1.
07-FEB-2006, entry version 8.
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Q7M504;
                                                                                                                                                                                                                                    Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (FEB-1993) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.
STRAIN=Balb/c; TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=94250690; PubMed=8193161;
Baumgartner B., Heiland S., Kunze N.,
"Conserved regulatory elements in the
promoters of mouse and man.";
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
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                                                                                                                                                                   GO:0008131; F:amine oxidase
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                                                                                                                                                                                           S71304;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            X70958; CAA50294.1; -; Genomic_DNA.
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1 AA;
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1240 MW;
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tive 0; Mismatc
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Pred. No.
                                                                                             21BEBFDCA4472DC7 CRC64;
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type I DNA topoisomerase
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                                                DB 2;
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K., Toyama
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H., Kumagai H.,
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RESULT 4
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DT 01-NOV-1996, i
DT 01-NOV-1996, e
DT 01-FEB-2006, e
DT 07-FEB-2006, e
DT 07-FEB-2006, e
COC REAGE PROTECTION ON NAME-REAGE;
OS Escherichia CC
CC Enterobacteria;
OX NCBL TAXID-56;
RN NUCLEOTIDE SE(
RX MEDLINE-91139;
RA TAO T., BOURNE
RX MEDLINE-91139;
RA TAO T., BOURNE
RX MEDLINE-9110;
CC Copyrighted b;
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                                                                                                                                RESULT 5
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ID Q50117 MY
AC Q50117 DT 01-NOV-11
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07-FEB-2006,
                                                                                                                                    EMBL; U19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Corynebacterineae; Mycobacteriaceae; NCBI_TaxID=1769;
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                                                                                                                                                                                                                                                                    Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                             Copyrighted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE Smith D.R.;
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Q50117;
                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (SEP-1994) to the EMBL/GenBank/DDBJ databases.
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Distributed under the Creative Commons Attribution-NoDerivs License
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J. Bacteriol. 173:1367-1375(1991).
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Score 22;
Pred. No.
                                                                                                                                    Genomic DNA.
CAABF1429DED5412
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Pred. No. le+04;
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Matches 5
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01-MAY-2000, sequence
07-FEB-2006, entry verse
Dopamine D5 (Fragment).
Gorilla gorilla (gorill
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-FEB-2006, entry version 6.
Fumarate hydratase (EC 4.2.1.2) (Fragment).
Fusarium sporotrichioides.
                                                                        NON_TER
                                                                                                                                                                                                                                                                                                              Q9TS39_9PRIM
Q9TS39;
                                                                                                        Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                                                                      "Transcription of a human dopamine D5 pseudogene."; Biochem. Biophys. Res. Commun. 181:16-21(1991).
                                                                                                                                                                                NUCLEOTIDE SEQUENCE.
MEDLINE=92068184; PubMed=1958184;
                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chow L.P., Fukaya N., Sugiura Y., Ueno Y., Submitted (OCT-1994) to the PIR data bank.
                                                                                                                                                              O' Dowd
                                                                                                                                                                       Nguyen
                                                                                                                                                                                                                                    Mammalia; Eutheria;
                                                               SEQUENCE
                                                                                                                                                                                                                NCBI_TaxID=9593;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GO:0004333; F:fumarate hydratase activity;
                                                                                                                                                             B. F.
                                                                                                                                                                       T., Sunahara R., Marchese A., Van Tol H.H.,
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33.3%;
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Pred. No.
                               Score 21.5;
Pred. No. 1.
                                                               39D8326BA63BB90B CRC64;
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RESULT 9
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05-JUL-2004,
07-FEB-2006,
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27-SEP-2005, sequence version 1.
07-MAR-2006, entry version 5.
60 kDa cell wall protein (Fragment).
Phaseolus vulgaris (Kidney bean) (French bean).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; endicotyledons;
rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae;
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             NON_TER
                                                                                                         NUCLEOTIDE SEQUENCE.

MEDLINE=98332731; PubMed=9666116; DOI=10.1016/S0378-1119(98)00265-0; Yin X.H., Gerbaud C., Francou F.X., Guerineau M., Virolle M.J.; Yin X.H., another amylolytic gene maps close to the amlB locus in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Robertson D., Mitchell G.P.,
 SEQUENCE
                                                 Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                      Streptomyces lividans TK24.";
Gene 215:171-180(1998).
                                                                                                                                                                         Streptomycineae;
NCBI_TaxID=1916;
                                                                                                                                                                                         Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
Streptomycineae; Streptomycetaceae; Streptomyces.
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-i- SUBCELLULAR LOCATION: Secreted; cell wall.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Slabas A.R.
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60 kDa cell wall protein.
N /FTId=PRO_0000079641.
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1160 MW;
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PubMed=9188482; DOI=10.1074/jbc.272.
chell G.P., Gilroy J.S., Gerrish C.,
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 F334C775A1A44871 CRC64;
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01-MAY-2000,
07-FEB-2006,
                                                                                                                                      NUCLEOTIDE SEQUENCE.
STRAIN-IOB 8413;
Grandvalet C., Coucheney F., Beltramo C., Guzzo J.;
"CtsR Is the Master Regulator of Stress Response Gene
                                                                                                                                                                                                                                                                                                           13-SEP-2005, integrated into I
13-SEP-2005, sequence version
07-FEB-2006, entry version 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MUCLEUILUB SECULION: NUMBELINE 92040090; PubMed=1718870; MEDLINE=92040090; PubMed=1718870; Peterson E.M., Cheng X., Markoff B.A., Fielder T.J., de la Maza L.M.; Functional and structural mapping of Chlamydia trachomatis species-reactific major outer membrane protein epitopes by use of neutralizing
                                                                                                                                                                                                                                                                                                                                                              Q4A3D9_OENOE
Q4A3D9;
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Q9R635;
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Distributed
                                                                                                                                                                                                                                            Bacteria; Firmicutes;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   monoclonal antibodies.";
Infect. Immun. 59:4147-4153(1991).
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Bacteria; Chlamydiae;
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                                                                                                                                                                                                                                                                                           ClpL2 protein (Fragment).
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                                                                                                         lnOenococcus oeni.";
J. Bacteriol. 487:5614-5623(2005).
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                      AJ890337;
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Lactobacillales; Oenococcus.
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01-OCT-1996, integrated into Unit
01-OCT-1996, sequence version 1.
07-FEB-2006, entry version 18.
Erythrocyte 65 kDa protein (P65)
                                                                                                                                                                                        PROTEIN SEQUENCE, AND CARBOHYDRATE-LINKAGE SITE MEDILINE=90004678; PubMed=2507249; MEDILINE=90004678; PubMed=2507249; MIDLINE=90004678; PubMed=2507249; MEDILINE=90004678; PubMed=2507249; MEDILINE=9004678; MEDILINE=90046788; MEDILINE=9004678; MEDILINE=9004678; MEDILINE=9004678; MEDILINE=9004678; MEDILINE=9004678; MEDILINE=9004678; MEDILINE=90046788; MEDILINE=9004678; MEDILINE=9004678; MEDILINE=9004678; MEDILINE=9004678; MEDILINE=90046788; MEDILINE=900467888; MEDILINE=900467888; MEDILINE=900467888; MEDILINE=900467888; M
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QSEDJ3;
15-MAR-2005, integrated into Unil
15-MAR-2005, sequence version 1.
07-FEB-2006, entry version 3.
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                                                           Copyrighted by the UniProt Consortium, see http://www.uniprot.org/termsDistributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Newton H.J., Hartland E.L.;
"Subtractive hybridization reveals DNA fragments preumophila, strain 02/41, and absent in Legionel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Legionella pneumophila.
Bacteria; Proteobacteria; Gammaproteobacteria; Legionellales;
Legionellaceae; Legionella.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; AY902880; AAW83806.1; NON_TER 1 1
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                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Distributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                                                                        a Found. Symp. 145:102-118(1989).
SUBCELLULAR LOCATION: Cytoplasm.
GO:0005737;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HUMAN
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C:cytoplasm; NAS
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SEQUENCE
  Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Meoptera; Endopterygota; Hymenoptera; Apocrita; Aculeata; Apo Andrenidae; Andreninae; Andrena; Callandrena. NCBI TaxID=205171;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gramene; Q9SB03; -.
GO; GO:0004373; F:glycogen (starch) synthase
GO; GO:0005982; P:starch metabolism; IEP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=98384837; PubMed=9718725; Hirano H., Eiguchi M., Sano Y.; Hirano H. Biguchi M., Sano Y.; This single base change altered the regulation of the Waxy gene at the post-cranscriptional level during domestication of rice."; Mol. Biol. Evol. 15:978-987(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oryza sativa (japonica cultivar-group).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; BEP clade;
Ehrhartoideae; Oryzeae; Oryza.
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Q9SB03;
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PubMed=16343953; DOI=10.1016/j.ympev.2005.10.003;

PubMed=16343953; DOI=10.1016/j.ympev.2005.10.003;

Larkin L.L., Neff J.L., Simpson B.B.;

"Phylogeny of the Callandrena subgenus of Andrena (Hymenoptera:

"Phylogeny of the Callandrena subgenus of Andrena (Hymenoptera:

"Phylogeny of the Callandrena subgenus of Andrena (Hymenoptera:

"Onvergent evolution.";

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                                                                                                                                                     Mitochondrion.
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Search completed: May 30, 2006, 11:09:59 Job time: 72.3333 secs

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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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| Active mi  | Human CD3 | Epitope w | Human COU | Human SCP | Active mi | hSARS vir | SARS viru | hSARS vir  |            | Hepatitis | Peptide i | F protein | Humanıze           | Humanized  | CDR3 trom | Anti-NKG2 | Anti-NKG2 | CD45RO/RB  | i. pylori   |          | •          |

### RESULT 1 ADR10602 Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody; Felis catus. Cat IgE epitope recognised 21-OCT-2004 ADR10602; ADR10602 standard; peptide; 15 15-JAN-2004; 2004WO-US003566 05-AUG-2004 WO2004065936-A2 (first entry) by monoclonal antibody 5.91, SEQ ID ₽

ALIGNMENTS

The present invention relates to a novel monoclonal antibody (I) that specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. (I) is useful for testing an allergen reactivity of an IgE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut and corn allergens. The sample is a biological sample collected from a dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal antibodies recognise epitopes on canine IgE corresponding to amino acid residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the

Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian Igor treating asthma or anaphylactic shock.

IgE

Example 6; Page 9; 14pp; English.

WPI; 2004-593545/57.

Hammerberg

16-JAN-2003; 2003US-0440472P

(UYNC-) UNIV NORTH CAROLINA STATE

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RESULT 2
ADR10601
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                                                                                                                                                                The present invention relates to a novel monoclonal antibody (I) that CC specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, eg. dog IgE. (I) is useful for testing an allergen reactivity of an IgE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut and corn allergens. The sample is a biological sample collected from a collec
                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian I or treating asthma or anaphylactic shock.
                                                                                                        Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 6; Page 9; 14pp; English.
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56.5%;
61.5%;
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Pred.
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48;
No.
DB 8;
0.24;
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                                 Length 15;
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Best Local Similarity

ADR10603

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RESULT 3
ADR10607
ID ADR1
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                                                                                                                                                                                                               The present invention relates to a novel monoclonal antibody (I) that CC specifically binds to a mammalian IgE epitope, where the epitope is compectation acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. CC between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. CC (I) is useful for testing an allergen reactivity of an IgE sample. The candidate corn allergens. The sample is a biological sample collected from a cc dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal cantibodies recognise epitopes on canine IgE corresponding to amino acid cresidues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the CC canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from CC cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and CC 3.76 were observed to have good cross-reactivity with the epsilon-chain cc of IgE from cat and horse, but did not exhibit cross-reactivity with ceither pig or human epsilon-chains of IgE. The present sequence is the corresponding or proposition of the construction of the corresponding to the corresponding to
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                                                                                    Query Match
Best Local S
Matches 6
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                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian IgE or treating asthma or anaphylactic shock.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pig IgE epitope recognised by monoclonal antibody 5.91,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-OCT-2004 (first entry
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                                                                                                           Similarity
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NDAPVQADRHSTT 14
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                                                                                      Conservative
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                                                                                                           45.9%;
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monoclonal antibody; 3.76 monoclonal antibody;
                                                                                      Score 39; DB
Pred. No. 9.9;
5; Mismatches
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RESULT 5
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ID ADC6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC specifically binds to a mammalian IgE epitope, where the epitope is CC between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. CC (I) is useful for testing an allergen reactivity of an IgE sample. The CC allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut CC and corn allergens. The sample is a biological sample collected from a CC dog, cat or horse. (I) is also useful for detecting mammalian IgE and for CC treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal CC residues recognise epitopes on canine IgE corresponding to amino acid CC residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the CC canine IgE spsilon-chain. Recognition of epsilon-chains from IgE from CC cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and CC of IgE from cat and horse, but did not exhibit cross-reactivity with CC either pig or human epsilon-chains of IgE. The present sequence is the CC horse IgE 5.91 recognition site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian I or treating asthma or anaphylactic shock.
Horse; immunoglobulin E; IgE; heavy chain; immunogen; allergy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15
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                                                     Horse immunoglobulin E, IgE, heavy chain immunogenic peptide P4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                present invention relates to a novel monoclonal antibody (I) that
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 NDSPVRTEQOATT 14
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23;
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                                                                                                                                                                                                                                                                                                                     Matches
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Best Local Similarity
                                                                                                                      AAU79709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunogenic composition comprising an isolated equine immunoglobulin E polypeptide that induces production of antibodies which specifically bind to equine immunoglobulin E.
                                                                                     15-JUL-2002
                                                                                                                                                       AAU79709 standard; peptide; 11 AA
                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15
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Pred. No. 52;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kalina WV;
                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                      7;
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                                                                                                                                                                                                                                                                                                                                                      Length 15
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                                                                                                                                                                                                                                                                                                                     Gaps
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Synthetic peptide CPP-2 used in invention of DRL90.

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Human; tissue-specific secretory polypeptide; DRL90; infection; cancer; immune disease; digestive disease; circulatory disease;

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RESULT 7
AAB38098
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           secretory polypeptide, DRL90, and the polynucrouser secretory polypeptide and encoding DNA are useful for diagno it. The DRL90 polypeptide and encoding DNA are useful for diagno treating cancer, immune diseases, infection, digestive diseases, circulatory diseases, endocrine diseases and infertility. The profit of the profi
                                                                                                                                                                                                                                                                             Human ABC1 cholesterol transporter; chromosome 9q31;
ATP-binding cassette; HHL deficiency disorder; high density lipoprotein;
Tangier disease; TD; familial HDL deficiency; FHA; polymorphism;
cardiovascular disease; coronary artery disease; coronary restenosis;
cerebrovascular disease; peripheral vascular disease;
Alzheimer's disease; Niemann-Pick disease; Huntington's disease;
X-linked adrenoleukodystrophy; cancer; gene therapy; genetic diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New tissue-specific secretory polypeptides applicable in diagnosis of remedies for cancer, immune diseases, infection, digestive diseases, circulatory diseases, endocrine diseases and infertility.
15-MAR-1999;
                                                 15-MAR-2000; 2000WO-IB000532
                                                                                                    21-SEP-2000.
                                                                                                                                                                                                                                                          prognosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB38098 standard; peptide; 10 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-SEP-2000; 2000JP-00293985.
29-SEP-2000; 2000JP-00302839.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-SEP-2001; 2001WO-JP008223
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   endocrine disease; infertility; cytostatic; immunomodulator;
antimicrobial; vasotropic; antiinfertility; hormonal; CPP-2.
                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ito Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200224908-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABC1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tanaka H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SPILTEKOA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SPVRTEQQA 12
                                                                                                                                                                                                                                                          prophylaxie;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FHA-3 mutant exon 41-encoded peptide fragment,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention relates to the isolation of a novel tissue-specific olypeptide, DRL90, and the polynucleotide sequence encoding 90 polypeptide and encoding DNA are useful for diagnosing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
99US-0124702P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nishimura A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     37.6%;
                                                                                                                                                                                                                                                          drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 32; DB 5;
Pred. No. 1.3e+02;
                                                                                                                                                                                                                                                          screening; transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 11;
                                                                                                                                                                                                                                                                animal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO:65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ne present
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CC a member of the ATP-binding cassette (ABC transporter) superfamily of intracellular cholesterol trafficking in monocytes and fibroblasts, being convolved in cholesterol efflux from the cell. The gene encoding ABC1 is clocated on chromosome 9g1, and mutations in this gene are associated with two genetic HBL (high density lipoprotein) deficiency disorders, compared disease (TD) and familial HBL deficiency (FHA). These diseases for an autosomal dominant trait. Low levels of HBL ("good colesterol") in the blood correlate with a high risk of cardiovascular codisease, coronary restenosis, and peripheral vascular disease. Conversely, a high level of HBL has protective effects against cardiovascular disease comprising the similate ABC1 actids, and methods of gene therapy for the treatment or prevention of corresponder encompasses compounds which mimic ABC1 activity, compounds which mimic abcoment of prevent cardiovascular disease, especially disease, coronary restenosis of gene therapy for the treatment or prevention also coresponds which mimic ABC1 activity, compounds which mimic abc1 disease due to polymorphisms in the CC further relates to methods for determining whether a patient has an correspond to the protess and nucleotides can be used to treat or corresponds. It for a standard disease, especially coronary artery disease, coronary restenosis or peripheral vascular disease, with the exact amino acid cardiovascular disease, especially coronary artery disease, especially excludes proteins with the exact amino acid cardio with the exact sequence as GenBank Accession No: AD12376. 1 The creator transporter peptide fragment used in the exemplifications of the invention
08-JUN-1999;
17-JUN-1999;
01-SEP-1999;
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to the human ABC1 cholesterol transporter protein (B38082) and to nucleic acid sequences (C69120) which encode it. ABC1 i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example; Fig 6E; 229pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New ABC1 polypeptide is useful for treating diseases associated with ABC1 biological activity, e.g. Alzheimer's disease, Huntington's disease and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (XENO-) XENON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYBR-) UNIV BRITISH COLUMBIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000-587528/55.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BIORESEARCH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0138048P.
99US-0139600P.
99US-0151977P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pimstone
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RESULT 8
AAB55476
ID AAB5
XX
AC AAB5
XX
AC AAB5
XX
AX
XX
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                                                                                                                                            Query Match
          07-MAR-2001
                                           AAB55476
                           AAB55476;
                                                                                                                                   Local
                                                                                                                         l Similarity
6; Conserv
                                           standard; peptide; 6 AA
                                                                                                         NDSPVRTEQQ 11
                                                                                        NDEDVRRERQ
                                                                                                                           Conservative
          (first entry)
                                                                                        10
                                                                                                                                  35.3%;
                                                                                                                          1;
                                                                                                                           Score 30; DB 3;
Pred. No. 2.6e+02;
1; Mismatches 3
                                                                                                                                            Length 10;
                                                                                                                            0,
                                                                                                                            Gaps
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0

Human elastase variant segment peptide SEQ ID NO:46

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AARS573
AARS573
ID AARS
XX AARS
AC AARS
XX 25-M
DT 16-N
XX dsDN
XX dsDN
XX dsDN
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XX dsDN
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XX Synt
XX Synt
PD 09-J
XX 09-J
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                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a purified elastase variant (I) with an amino acid sequence different from that of a precursor elastase, the difference comprising a substitution of an active site histidine residue corresponding to residue 43 in human neutrophil elastase with a different amino acid residue so that (I) has substrate specificity substantially different from the precursor elastase. (I) has cytostatic activity, and can be used in antibody-directed enzyme activated prodrug therapy. The elastase variant can be used to cleave a particular substrate, especially those containing histidine residues at the substrate site. Site-specific proteolysis is useful in therapeutic at the substrate site. Site-specific proteolysis is useful in therapeutic at phications, e.g. for antibody-directed enzyme activated prodrug therapy (ADEPT). AAC88022, AAC88023 and AAB55532 to AAB55526 represent sequences used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; elastase; human neutrophil
                                                                                                      dsDNA-dependent kinase inhibitor; fatty acyl-peptide; conjugate; antiproliferative; tumor; psoriasis; docosahexaenoic acid; DHA; eicosapentaenoic acid; EPA; antitumor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Carter PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-MAY-2000; 2000WO-US006692.
                                                                                                                                                                   dsDNA-dependent kinase inhibitor
                                                                                                                                                                                               25-MAR-2003
16-NOV-1994
                                                                                                                                                                                                                                              AAR55733;
                                                                                                                                                                                                                                                                         AAR55733 standard; peptide; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 4; Fig 3; 79pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antibody-directed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Elastase variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antibody-directed enzyme
               09-JUN-1994
                                              WO9412530-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention
                                                                                                                                                                                                                                                                                                                                                                                  10
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                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                  QQATTW 15
                                                                                                                                                                                                                                                                                                                                                      QQHTTW 6
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                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                               (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0132640P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (H43A) having altered substrate specificity useful denzyme activated prodrug therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  variant; substrate; mutant; mutagenesis; histidine; elastase; H43A; cytostatic; proteolysis; ADEPT; d enzyme activated prodrug therapy.
                                                                                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                 34.1%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rodriques
                                                                                                                                                                                                                                                                                                                                                                                                                 Score 29; DB Pred. No. 2.1e
                                                                                                                                                                                                                                                                                                                                                                                                                  <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                  .1e+06
                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 6;
                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiinflammatory; cysteine protease Harada's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The peptides given in AAR55718-48 can each be conjugated through an amide linkage with a polyunsaturated fatty acid moinery, such as docosahexaenoic acid or eicosapentanoic acid, to improve antiproliferative activity. The dsDNA-dependent kinase inhibitor given in AAR55733 competes with native kinases associated with meoplastic cell proliferation or transformation, psoriasis, etc. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New fatty acyl-peptide conjugates for inhibiting cell proliferation more active than free peptide, partic. for treating tumours, virus-
                                                                                                                       Novel cysteine protease inhibitory protein, u
for diagnosing Behcet's disease and Harada's
                                                                                                                                                                                                                                                                                                                                                                          05-MAR-2003; 2003JP-00059082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cysteine protease inhibitory protein fragment #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-DEC-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADR97089
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig 1; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1994-200194/24.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-NOV-1992;
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                                                                                                                                                                                                           WPI; 2004-668624/65.
                                                                                                                                                                                                                                                                                                                                                                                                                            05-MAR-2004; 2004WO-JP002902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO2004078975-A1
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                                                                                                                                                                                                                                                                                                                     (SENP ) SENJU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   YNDNPMEEEE 15
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                                                                        SEQ ID NO 4; 64pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          34.18; ilarity 40.08; Conservative
                                                                                                                                                                                                                                                                Shiota H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry
                                                                                                                                                                                                                                                                                                                     PHARM CO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KUTATO FEJLESZTO KFT PEPTIDES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vasotropic; immunost
inhibitory protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 29; DB
Pred. No. 6.1e
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunostimulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Szederkenyi F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis; Behcet's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6.1e+02;
ches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 2;
                                                                                                                                  useful as diagnostic marker 
s disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Horvath
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Balogh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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A cysteine protease inhibitory ADR97086 at its N-terminal and

protein (I) having the amino acid sequence exhibiting the molecular weight of 31 kDa,

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RESULT 11
AAMS6548
ID AAMS6
XX AAMS6
XX DE TOXIN
XX Photo
KW Photo
KW Ucb;
KW Ucb;
KW Ucb;
KW Uweste
KW Corn
XX Photo
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       á
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          អ្នក្សក្សដុឌ្ធ
ស្រុក្សស្ត្រដុឌ្ធ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                      They can be used to control insects pests from the Lepidoptera, Coleoptera, Hymenoptera, Diptera, Dictyoptera, Acarina or Homoptera orders, especially the Southern or Western corn rootworm, Colorado porders, especially the Southern or Western corn rootworm,
                                                                                                   This is a symbiotic bacterium of the nematodes of the Heterorhabditis genus. The bacterium has at least 4 distinct genomic regions, tca, tcb, tcc, and tcd. Peptide products are produced from these regions that are associated with insecticidal activity. The native toxins are secreted proteins. The proteins are toxic to insects upon exposure and especially when ingested. The nucleic acid sequence can be used to produce transgenic plants, baculoviruses or microbial hosts for toxin production.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               is new. (I) is useful for screening a compound or its salt having activity of promoting or inhibiting the function of (I). (I) enables diagnosis, prevention or treatment of Behcet's disease or Harada's disease. This sequence is a fragment of the cysteine protease inhibitory
                                                                                                                                                                                                                                                                                            The present sequence represents a fragment of the toxic protein TcaC (encoded by gene tcaC) of the bacterium Photorhabdus luminescens (W-14).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (DOWC )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tcb; tcc; tcd; insecticidal activity; toxin; Lepidoptera; Coleoptera; Hymenoptera; Diptera; Dictyoptera; Acarina; Homoptera; Southern; Western corn rootworm; Colorado potato beetle; mealworm; boll weevil;
                                                                                                                                                                                                                                                                                                                                                                                                                              Isolated toxins from Photorhabdus luminescens strains -control of insect pests.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1998-179427/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-AUG-1996;
06-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ffrench-Constant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Photorhabdus luminescens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Photorhabdus luminescens W-14; nematode; symbiotic; Heterorhabditis; tca;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW56548 standard; peptide; 12 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW56548
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                grub; beetle armyworm;
                                                                                                                                                                                                                                                                                                                                                                          30; Page 151; 321pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'n,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DOWELANCO.
WISCONSIN ALUMNI RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Constant RH,
Roberts JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DSPVRTEQQATTW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DRPARHPQEQPLW 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ą
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bowen DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96US-00705484.
96US-00743699.
96WO-US018003.
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     1109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          34.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rocheleau TA,
Strickland JA,
     weevil,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          corn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Petell J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          borer; Tobacco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                black cutworm;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bacterium Photorhabdus luminescens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TA, Blackburn MB, H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          No.
  grub,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6.1e+02
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     beetle armyworm,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cabbage looper; codling moth;
hornworm; budworm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Schoonover S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hey
TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TD, Merlo I
Sukhapinda
                           Colorado potato
     black cutworm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , D.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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Best Loc Matches Query Match Best Local

Similarity 4; Conserv

Conservative

32.9%;

Score 28; DB Pred. No. 2.1e 2; Mismatches

<u>,</u>

0

Gaps

0

DB 2; .1e+06;

Length 9;

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RESULT 12
AM76034
ID AM76034
AC AAM76
AC AAM76
AC AAM76
AC AM760
AC LM609
AC UM609
AC AAM70
ACC AAM70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                            AAW76007-W76040 are protein fragments of the grafted monoclonal antibody LM609 heavy and light chain variable region. LM609 and the antibody vitaxin bind selectively to integrin alphavbeta3 and can be used to inhibit binding of alphavbeta3 to a ligand and thus block integrin-mediated signal transduction. This is useful in the treatment, prevention and diagnosis of alphavbeta3-mediated disease, specifically angiogenesis and restenosis (but also e.g. (non-)lmmune inflammation, diabetic retinopathy, neovascular glaucoma, cancer, psociasis, rheumatoid arthritis, macular degeneration, osteoporosis etc.). The antibodies contain non-murine framework regions so are suitable for use in humans. Enhanced types of LM609 have affinity more than 90 times greater than
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cabbage
tobacco
  Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Humanised antibody, Vitaxin, that binds selectively to alphaVbeta3 integrin - and related grafted antibodies based on murine monoclon LM609, also related nucleic acid, used to treat, prevent or diagno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LM609; inhibitor; integrin-mediated signal transduction; treatment; diagnosis; angiogenesis; restenosis; inflammation; diabetic retinopathy; nevvascular glaucoma; cancer; psoriasis; rheumatoid arthritis; murine; macular degeneration; osteoporosis; primer; V-L region; CDR;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 62;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       angiogenesis or restenosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Huse WD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9833919-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  complementarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LM609 grafted antibody V-L region CDR3 protein fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-NOV-1998 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (IXSY-) IXSYS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1998-437472/37
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                                                          parent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV49871.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             looper, codling moth, hornworm or budworm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DSPVRTEQOATTW 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antibody; variable region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DSP---EVSITTW 12
                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Glaser SM
     Ą
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                                                             the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               determining
                                                       parent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     33.5%;
                                                             antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                  English
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Pred. No. 5.8e+02;
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RESULT 14
ABO19830
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AC ABO19
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AC ABO19
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AAB61392
JID AAB61392
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AC AAA
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to enhanced LM609 grafted antibodies exhibiting selective binding affinity to alphaVbeta 3 integrin or their functional fragments. The antibodies or their functional fragments can be used in the diagnosis and treatment of alphaVbeta 3-mediated diseases such as angiogenesis, inflammatory diseases (such as psoriasis and chronic articular rheumatism), disorders associated with inappropriate or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Enhanced LM609 grafted antibodies exhibiting selective binding affinity to alpha(V)beta(3) integrin, useful in the diagnosis and treatment of angiogenesis, inflammatory diseases e.g. psoriasis, cancers and osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inopportune invasion of vessels (such as diabetic retinopathy, neovascular glaucoma and cancer disorders such as tumours and sarcoma), retinal diseases (such as macular degeneration), res
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LM609; grafted antibody; alphaVbeta_3 integrin; angiogenesis;
inflammatory; cancer; retina; restenosis; osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 41; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-050110/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             osteoporosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Huse WD, Wu H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified
  Enhanced LM609 light chain variable region CDR3
                                                          28-AUG-2003
                                                                                                                                                                ABO19830 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MOLE-) APPLIED MOLECULAR EVOLUTION
                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
mes 4; Conserv
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                                                          entry)
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Pred. No. 2.1e
2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 4;
2.1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 uch as tumours and Kaposi's degeneration), restenosis
                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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RESULT 15
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Best Local
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alpha_vbeta_3-mediated disease; complementarity determining region;
restenosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a Vitaxin or LM609 grafted antibody, exhibiting selective binding affinity to alpha vbeta 3. The Vitaxin or LM609 grafted antibody is useful for treating an alpha-vbeta_3-mediated disease e.g. angiogenesis or restenosis. The present sequence represents the amino acid sequence of a LM609 complementarity determining region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New Vitaxin or LM609 grafted antibody exhibiting selective binding affinity to alphavbeta3, useful for treating an alphavbeta3-mediated disease e.g., angiogenesis or restenosis.
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Huse WD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-JAN-1998;
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                                                                                                                                                                                                           Surface simulation synthetic peptide useful in HIV-1 vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 65; Page 12; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB;
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                                                     01-DEC-2001; 2001US-00012806
                                                                                                 12-JUN-2003
                                                                                                                                           Synthetic
                                                                                                                                                       Human immunodeficiency virus
                                                                                                                                                                           virucide; epitope.
                                                                                                                                                                                        Surface
                                                                                                                                                                                                                                   22-SEP-2003
                                                                                                                                                                                                                                                        ABR62303;
                                                                                                                                                                                                                                                                             ABR62303 standard; peptide; 9
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N-PSDB; ACD30189.
                                                                          23-NOV-2002; 2002WO-US037664
                                                                                                                      WO2003048186-A2
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                                (CREV/) CREVECOEUR
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                                                                                                                                                                                      simulation
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                                                                                                                                                                                      synthetic
                                                                                                                                                                                                                                                                                                                                                                                          32.9%;
                                                                                                                                                                                                                                                                                                                                                                               Score 28; DB 6;
Pred. No. 2.1e+06;
2; Mismatches (
                                                                                                                                                                                       peptide;
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                                                                                                                                                                                         gssp;
                                                                                                                                                                                         HIV-1; vaccine; anti-HIV;
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Crevecoeur H;

WPI; 2003-558941/52.

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Search completed: May 30, 2006, 10:12:40 Job time : 62 secs
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                                                                                                                                                                 Query Match 32.9%;
Best Local Similarity 62.5%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                          The present sequence is that of a surface simulation synthetic peptide (SSSP) corresponding to a non-sequential conserved residue epitope of HIV -1 glycoprotein gp120 required for viral pathogenicity. It is an example of SSSPs of the invention, which incorporate amino acid sequences that simulate the 3-dimensional spatial positions of non-sequential conserved residues necessary for viral pathogenicity. The SSSPs are useful for incorporation into vaccines effective in eliciting an effective broad spectrum immune response against hyper-variable viral pathogens such as HIV-1, and in diagnostic kits. SSSPs provide a reliable strategy for allowing the immune system to process and recognise discontinuous epitopes and to mount immune responses to the 3-dimensional configurations of targeted proteins
                                                                                                                                                                                                                                                        Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Configuration of synthetic peptides useful for treating hyper-variable viral pathogen e.g. HIV, involves identifying non-sequential conserved residues in selected viral proteins and designing a sequence from image scan of the residues.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 27; 55pp; English.
                                                                                                                        4 SPVRTEQQ 11
                                                                                  NPCRTEKO 8
                                                                                                                                                                 Score 28; DB 6; Pred. No. 2.1e+06; 2; Mismatches 1
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Minimum
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Listing first 45 summaries
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length: 15
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Match
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2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

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US-10-381-089-18

US-10-381-089-18

US-10-447-465-65

US-10-447-34-65

US-10-147-34-65

US-10-148-2794A-2

US-10-262-794A-2

US-10-262-794A-2

US-10-305-231-08

US-10-463-847-88

US-10-467-546-31

US-10-296-656-31

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US-11-126-738-12
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10 Appli

11 Appli

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13 Appli

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15 Appli

16 Appli

17 Appli

18 Appli
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|             | 45                | 44                | 43                | 42                | 41                | 40                | 39                | 38                | 37                | 36                | 35                | 34                | 33                 | 32                 | 31                 | 30                | 29                | 28                |
|-------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|--------------------|--------------------|--------------------|-------------------|-------------------|-------------------|
|             | 26                | 26                | 26                | 26                | 26                | 26                | 26                | 26                | 26                | 26                | 26                | 26                | 27                 | 27                 | 27                 | 27                | 27                | 27                |
|             | 30.6              | 30.6              | 30.6              | 30.6              | 30.6              | 30.6              | 30.6              | 30.6              | 30.6              | 30.6              | 30.6              | 30.6              | 31.8               | 31.8               | 31.8               | 31.8              | 31.8              | 31.8              |
|             | 10                | 10                | 10                | 10                | 10                | 9                 | 9                 | 9                 | 9                 | 9                 | 9                 | 9                 | 14                 | 14                 | 14                 | 13                | 13                | 13                |
|             | ហ                 | υ                 | 4                 | w                 | w                 | σ                 | σ                 | σ                 | 6                 | v                 | 4                 | 4                 | σ                  | v                  | σ                  | w                 | w                 | ω                 |
| AT.TGNMENTS | US-10-929-988-436 | US-10-659-207-436 | US-10-117-937-561 | US-09-783-931-53  | US-09-908-322-53  | US-11-203-251A-75 | US-11-203-251A-10 | US-11-067-159-560 | US-11-067-064-560 | US-10-883-020-19  | US-10-428-335-103 | US-10-117-937-560 | US-11-004-399-2037 | US-10-807-807-1344 | US-10-808-187-1344 | US-09-873-224-244 | US-09-878-381-244 | US-09-899-046-244 |
|             | Sequence 436, App | Sequence 436, App | Sequence 561, App | Sequence 53, Appl | Sequence 53, Appl | Sequence 75, Appl | Sequence 10, Appl | Sequence 560, App | Sequence 560, App | Sequence 19, Appl | Sequence 103, App | Sequence 560, App | Sequence 2037, Ap  | Sequence 1344, Ap  | Sequence 1344, Ap  | Sequence 244, App | Sequence 244, App | Sequence 244, App |

#### Sequence 2, Application US/10758165 ; Publication No. US20050196816A1 ; GENERAL INFORMATION: APPLICANT: Hammerberg, Bruce TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES FILE REFERENCE: 5051-661 ; CURRENT APPLICATION NUMBER: US/10/758,165 ; CURRENT FILING DATE: 2004-01-16 ; PRIOR APPLICATION NUMBER: US 60/440,472 ; PRIOR FILING DATE: 2003-01-16 ; NUMBER OF SEQ ID NOS: 16 ; SOFTWARE: Patentin version 3.2 ; SEQ ID NO 2 ; LENGTH: 15 밁 Ś US-10-758-165-1 RESULT 2 US-10-758-165-1 ; TYPE: PRT; ORGANISM: Felis catus US-10-758-165-2 US-10-758-165-2 RESULT 1 Sequence 1, Application US/10758165 Publication No. US20050196816A1 GENERAL INFORMATION: APPLICANT: Hammerberg, Bruce TITLE OF INVENTION: IMMUNOSLOBULIN E DETECTION IN MAMMALIAN SPECIES FILE REFERENCE: 5051-661 CURRENT APPLICATION NUMBER: US/10/758,165 CURRENT FILING DATE: 2004-01-16 PRIOR APPLICATION NUMBER: US 60/440,472 PRIOR FILING DATE: 2003-01-16 NUMBER OF SEQ ID NOS: 16 SOFTWARE: PatentIn version 3.2 SEQ ID NO 1 Matches Query Match Best Local Similarity LENGTH: 15 TYPE: PRT ORGANISM: Canis familiaris 15; Conservative 1 HNDSPVRTEQQATTW 15 1 HNDSPVRTEQQATTW 15 100.0%; 0, Score 85; DB 5; Pred. No. 7e-08; Mismatches ۍ ۲: 0 Length 15; Indels 0 Gaps

0

Query Match

56.5%;

Score 48;

DB 5;

Length 15;

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Sequence 3, Application US/10758165
Publication No. US20050196816A1
GENERAL INFORMATION:
APPLICANT: Hammerberg, Bruce
ITITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-61
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR RILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 16
SOPTWARE: Patentin version 3.2
SEQ ID NO 3
LENGTH: 15
TYPE: PRT
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US-10-052-788-4
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US-10-758-165-3
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US-10-758-165-3
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TYPE: PRT
ORGANISM: Sus scrofa
US-10-758-165-7
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US-10-758-165-7
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Sequence 4, Application US/10052788
Publication No. US20030087314A1
GENERAL INFORMATION:
APPLICANT: Gershwin, Laurel J.
APPLICANT: Pettigrew, Howard David
                                                                                                                                                                                                                 Query Match
Best Local S
Matches 7
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TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MARMALIAN SPECIES
FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR APPLICATION ORDER: US 60/440,472
PRIOR PILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PATENTIN VERSION 3.2
SEQ ID NO 7
SEQ ID NO 7
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Publication No. US20050196816A1
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 Howard David
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Pred. No. 10;
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APPLICANT: Kalina, Warren V.

APPLICANT: The Regents of the University of California
FITLE OF INVENTION: Epsilon Immunoglobulin Chain Derived Peptides for
FITLE OF INVENTION: Induction of Anti-IgE Antibodies
FILE REFERENCE: 023070-121000US
CURRENT APPLICATION NUMBER: US/10/052,788
CURRENT FILING DATE: 2001-11-08
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/381,089
CURRENT FILING DATE: 2003-03-20
PRIOR APPLICATION NUMBER: JP 2000-293985
PRIOR FILING DATE: 2000-09-22
PRIOR APPLICATION NUMBER: JP 2000-302839
PRIOR FILING DATE: 2000-09-29
NUMBER OF SEQ ID NOS: 23
SEQ ID NO 18
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                                                                                                                                    US-10-381-089-8
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GENERAL INFORMATION:
APPLICANT: ITO, Yasuaki
APPLICANT: TANAKA, Hideyuki
APPLICANT: NISHIMURA, Atsushi
APPLICANT: OGI, Kazuhiro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18, Application US/10381089 Publication No. US20040053276A1 GENERAL INFORMATION:
                                                                                               Sequence 8, Application US/10381089 Publication No. US20040053276A1
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Best Local
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Best Local :
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APPLICANT: TANAKA, Hideyuki
APPLICANT: NISHIMURA, Atsus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: NISHIMURA, Atsushi
APPLICANT: OGI, Kazuhiro
TITLE OF INVENTION: Tissue specific novel secretory polypeptide and its DNA
FILE REFERENCE: 2789 uSOP
                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
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66.7%;
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Pred. No.
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Pred. No. 1.2e+02;
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TITLE

OF INVENTION: Tissue specific novel secretory polypeptide and its

DNA

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; LENGTH: 15
; TYPE: PRT
; ORGANISM: Human
US-10-381-089-8
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US-10-617-334-65
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CURRENT FILING DATE: 2003-06-02
PRIOR APPLICATION NUMBER: US 09/526,193
PRIOR FILING DATE: 2000-03-15
PRIOR APPLICATION NUMBER: 60/124,702
PRIOR APPLICATION NUMBER: 60/124,702
PRIOR FILING DATE: 1999-06-08
PRIOR FILING DATE: 1999-06-08
PRIOR FILING DATE: 1999-06-08
PRIOR APPLICATION NUMBER: 60/139,600
PRIOR FILING DATE: 1999-06-07
PRIOR APPLICATION NUMBER: 60/151,977
PRIOR APPLICATION NUMBER: 60/151,977
PRIOR APPLICATION NUMBER: 60/151,977
PRIOR FILING DATE: 1999-09-01
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US-10-452-510-65
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US-10-452-510-65
                                                             Sequence 65, Application US/10617334
Publication No. US20040058869A1
GENERAL INFORMATION:
APPLICANT: Hayden, Michael R.
APPLICANT: Brooks-Wilson, Angela R.
APPLICANT: Brooks-Wilson, Angela R.
TITLE OF INVENTION: METHODS AND REAGENTS FOR MODULATING CHOLESTEROL LEVELS
FILE REFERENCE: 760050-91
                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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LENGTH: 10
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Best Local Similarity 66.7%;
Matches 6; Conservative
                      CURRENT APPLICATION NUMBER: US/10/617,334
CURRENT FILING DATE: 2003-07-10
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SOFTWARE: FastSEQ for Windows Version
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                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity 60.0%; les 6; Conservation
APPLICATION NUMBER: US 09/526,193
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Pred. No. 2.5e+02;
1; Mismatches
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Pred. No. 1.7e+02;
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PRIOR FILING DATE: 2000-03-15
PRIOR APPLICATION NUMBER: 60/124,702
PRIOR FILING DATE: 1999-03-15
PRIOR FILING DATE: 1999-03-15
PRIOR FILING DATE: 1999-06-08
PRIOR FILING DATE: 1999-06-08
PRIOR REPLICATION NUMBER: 60/139,600
PRIOR TILING DATE: 1999-06-17
PRIOR APPLICATION NUMBER: 60/151,977
PRIOR APPLICATION NUMBER: 60/151,977
PRIOR APPLICATION NUMBER: 50/151,977
PRIOR FILING DATE: 1999-09-01
NUMBER OF SEQ ID NOS: 287
SOFTWARE: PATENTIN 3.0
SEQ ID NO 65
LENGTH: 10
                                                                                                                                                   Query Match
Best Local Similarity
Thes 6; Conserve
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Best Local Similarity
Watches 6; Conserve
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US-10-833-679-65
                                                                                                                                                                                                                                                               ; ORGANISM: Homo sapiens
US-10-744-465-65
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CURRENT FILING DATE: 2003-12-23
PRIOR APPLICATION NUMBER: 10/617,334
PRIOR HILING DATE: 2003-07-10
PRIOR APPLICATION NUMBER: US 09/526,193
PRIOR FILING DATE: 2000-03-15
PRIOR APPLICATION NUMBER: 06/124,702
PRIOR APPLICATION NUMBER: 60/124,702
PRIOR PRILING DATE: 1999-03-15
PRIOR APPLICATION NUMBER: 60/138,048
PRIOR FILING DATE: 1999-06-08
PRIOR FILING DATE: 1999-06-08
PRIOR FILING DATE: 1999-06-08
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                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1999-06-17
PRIOR APPLICATION NUMBER: 60/151,977
PRIOR FILING DATE: 1999-09-01
NUMBER OF SEQ ID NOS: 287
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 65
LENGTH: 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Hayden, Michael R. APPLICANT: Brooks-Wilson, An APPLICANT: Pimstone, Simon N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Brooks-wilson, Angela R.
APPLICANT: Pimstone, Simon N.
TITLE OF INVENTION: METHODS AND REAGENTS FOR MODULATING CHOLESTEROL LEVELS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/139,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                 NDSPVRTEQQ 11
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                                                                                  NDEDVRRERQ 10
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Pred. No. 2.5e+02;
1; Mismatches 3
                                                                                                                                                                                                 Score 30; DB 4;
Pred. No. 2.5e+02;
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Sequence 65, Application US/10833679
Publication No. US20040185508A1
GENERAL INFORMATION:

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Query Match
Best Local Similarity
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LENGTH: 10
TYPE: PRT
  Query Match
Best Local Similarity
                                                                                                      SOFTWARE: PatentIn 3.0
SEQ ID NO 65
LENGTH: 10
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 65, Application US/10818279 Publication No. US20050136421A1 GENERAL INFORMATION:
                                                              ORGANISM: Homo sapiens -10-818-279-65
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APPLICANT: Brooks-Wilson, Angela R.
APPLICANT: Pimstone, Simon N.
TITLE OF INVENTION: METHODS AND REAGENTS FOR MODULATING CHOLESTEROL LEVELS
FILE REFERENCE: 760050-126
                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 10/745,377
PRIOR FILING DATE: 2003-12-23
PRIOR APPLICATION NUMBER: 10/617,334
PRIOR FILING DATE: 2003-07-10
PRIOR APPLICATION NUMBER: 09/526,193
PRIOR FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/818,279
CURRENT FILING DATE: 2004-04-05
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SOFTWARE: PatentIn 3.0
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PRIOR FILING DATE: 2003-06-02
PRIOR APPLICATION NUMBER: 10/617,334
PRIOR FILING DATE: 2003-07-10
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CURRENT APPLICATION NUMBER: US/10/833,679
CURRENT FILING DATE: 2004-04-28
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TITLE OF INVENTION: METHODS AND REAGENTS FOR MODULATING CHOLESTEROL LEVELS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OR APPLICATION NUMBER: 09/526,193
OR FILING DATE: 2000-03.15
OR APPLICATION NUMBER: 60/124,702
OR FILING DATE: 1999-03-15
OR APPLICATION NUMBER: 60/138,048
OR FILING DATE: 1999-06-08
                                                                                                                                                                                                                                               FILING DATE: 1999-06-08
APPLICATION NUMBER: 60/
FILING DATE: 1999-06-17
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60.0%;
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Pred. No. 2.5e+02
  Score
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  30;
  DB 5; ; 2.5e+02;
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                     Length 10;
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                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/395,497
FILING DATE: 28-FEB-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/007,255
FILING DATE: 06-NOV-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/608,423
FILING DATE: 28-FEB-1996
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/705,484
PRILORION NUMBER: US 08/705,484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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APPLICATION NUMBER: US 08/851,567
FILING DATE: 05-MAY-1997
CLASSIFICATION:
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APPLICATION NUMBER: US 0
FILING DATE: 18-MAY-1993
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                      TELECOMMUNICATION INFORMATION: TELEPHONE: 608-251-5000
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CITY: Indianapolis
TELEPHONE: 6
                                                                       REFERENCE/DOCKET NUMBER:
                                                                                             NAME: Seay, Nicholas REGISTRATION NUMBER:
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9330 Zionsville Road
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Bowen, David J
Petell, James
Fatig, Raymond
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608-251-9166
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Sukhapinda, Kitisri
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Blackburn, Michael B.
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, Nicholas J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US 08/063,615
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Query Match
Best Local Similarity
Thes 7; Conserve
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; Sequence 88, Application US/09900590; Publication No. US20030028009A1; GENERAL INFORMATION: Usiliam D
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                                                                                 Query Match
Best Local (
                                                                   Matches
                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 88: SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION UMBER: US/09/900,590
FILING DATE: 06-Jul-2001
CLASSIFICATION: CUNknown>
                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
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                                                                                                                                                                                                                                                                                          NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-IX 2965
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 88:
                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/016,061
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Huse, William D.
APPLICANT: Huse, William D.
TITLE OF INVENTION: Anti-Alpha V Beta 3 Recombinant Human
Antibodies, Nucleic Acids Encoding Same and Methods of Use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
                                10 QQATTW 15
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amino acid
                                                                   4; Conservative
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QQSTSW 6
                                                                                                                                                                                                      LENGTH: 9 amino acids
TYPE: amino acid
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STATE: California
                                                                                                                                                                                                                                                                             TELEFAX:
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ilarity 53.8%;
Conservative
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                                                                                   32.9%;
                                                                 Score 28; DB 3; Le
Pred. No. 1.9e+06;
2; Mismatches 0;
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                                                                                                    Length 9;
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RESULT 15

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US-10-012-806A-39

Sequence 39, Application US/10012806A

Publication No. US20030125518A1

GENERAL INFORMATION:

APPLICANT: CREVICOUER, HARRY

INTILE OF INVENTION: SURFACE SIMULATION SYNTHETIC PEPTIDES USEFUL IN THE

TITLE OF INVENTION: TREATMENT OF HYPER-VARIABLE VIRAL PATHOGENS

FILE REFERENCE: 2001-Crev1

CURRENT APPLICATION NUMBER: US/10/012,806A

CURRENT FILING DATE: 2002-03-12

NUMBER OF SEQ ID NOS: 45

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 39
                                                                                                                                                                                                                                          ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: peptide US-10-012-806A-39
Search completed: May 30, 2006, 10:24:41 Job time : 48 secs
                                                                                  В
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                                                                                                                                                                  Matches
                                                                                                                                                                                                      Query Match
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TYPE: PRT
                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                       32.9%;
Local Similarity 62.5%;
nes 5; Conservative
                                                                                                                        4 SPVRTEQQ 11
                                                                                  NPCRTEKO 8
                                                                                                                                                                Score 28; DB 4; Le
Pred. No. 1.9e+06;
2; Mismatches 1;
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Maximum Match 100%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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length: 15
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US-11-297-317-21

US-11-127-886-778

US-11-122-986-778

US-10-489-071-6

US-10-538-066-163

US-10-546-594-92

US-11-140-487A-1766

US-10-516-079-64

US-11-10-981-300-34

US-11-10-981-300-34

US-11-106-014-84

US-11-106-014-84

US-11-106-014-84

US-11-106-014-84

US-11-10538-066-159

US-10-538-066-159

US-11-023-959A-18

US-11-023-959A-18

US-11-023-959A-18

US-11-023-959A-18

US-11-023-959A-18

US-11-023-959A-72

US-11-33-056-184

US-10-538-066-268

US-10-538-066-413

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21, Appl
778, App
163, Appl
192, Appli
92, Appli
94, Appl
34, Appl
36, Appl
37, Appl
18, Appl
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| 45                  | 44                  | 43                  | 42                | 41                | 40               | 39                  | 38                  | 37                  | 36                  | 35                 | 34                | ω<br>ω            | 32               | 31              | 30               | 29                  | 28                | 27                |   |
|---------------------|---------------------|---------------------|-------------------|-------------------|------------------|---------------------|---------------------|---------------------|---------------------|--------------------|-------------------|-------------------|------------------|-----------------|------------------|---------------------|-------------------|-------------------|---|
| 18                  | 18                  | 18                  | 18                | 18                | 18               | 18                  | 18                  | 18                  | 18                  | 18                 | 18                | 18                | 18               | 18              | 18               | 19                  | 19                | 19                |   |
| 21.2                | 21.2                | 21.2                | 21.2              | 21.2              | 21.2             | 21.2                | •                   | •                   | 21.2                |                    | 21.2              | 21.2              | 21.2             |                 | 21.2             | 22.4                | 22.4              | 22.4              |   |
| 10                  | 10                  | 10                  | 10                | 10                | 10               | 9                   | 9                   | 9                   | 9                   | 9                  | 9                 | 9                 | œ                | 7               | σ                | 15                  | 13                | 12                |   |
| 7                   | 7                   | 7                   | σ                 | σ                 | σ                | 7                   | 7                   | 7                   | 7                   | 7                  | σ                 | σ                 | 7                | σ               | σ                | 7                   | 7                 | σ                 |   |
| US-11-140-487A-1869 | US-11-140-487A-1768 | US-11-140-487A-1378 | US-10-538-066-267 | US-10-538-066-172 | US-10-499-266-20 | US-11-140-487A-1767 | US-11-140-487A-1561 | US-11-140-487A-1377 | US-11-140-487A-1042 | US-11-140-487A-895 | US-10-538-066-299 | US-10-538-066-174 | US-11-213-668-74 | US-10-522-043-2 | US-10-525-126-33 | US-11-140-487A-2231 | US-11-121-282-197 | US-10-538-066-381 |   |
| Seguence            | Sequence            | Sequence            | Sequence          | Sequence          | Sequence         | Sequence            | Sequence            | Sequence            | Sequence            | Sequence           | Sequence          | Sequence          | Sequence         | Sequence        | Sequence         | Sequence            | Sequence          | Sequence          | , |
| 1869, Ap            | 1768, Ap            | 1378, Ap            | 267, App          | 172, App          | 20, App1         | 1767, Ap            | 1561, Ap            | 1377, Ap            | 1042, Ap            | 895, App           | 299, App          | •                 | 74, App1         | 2, Appli        | 33, App1         | 2231, Ap            |                   | 381, App          |   |

# ALIGNMENTS

US-11-297-317-21

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RESULT 2
US-11-140-487A-1220
US-21-140-487A-1220
; Sequence 1220, Application US/11140487A
; Publication No. US20060093617A1
; GENERAL INFORMATION:
; APPLICANT: Innogenetics N.V.
; TITLE OF INVENTION: Peptides for inducing a CTI
; FILE REFERENCE: 166
; CURRENT APPLICATION NUMBER: US/11/140,487A
; CURRENT FILING DATE: 2005-05-31
; PRIOR APPLICATION NUMBER: EP 04012951.2
; PRIOR FILING DATE: 2004-06-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Gregorio Aversa
APPLICANT: Gregorio Aversa
APPLICANT: Gregorio Aversa
APPLICANT: Jose M. Carballido Herrera
APPLICANT: Jose M. Carballido Herrera
APPLICANT: Jose M. Saldanha
APPLICANT: Jose M. Hall
ITITLE OF INVENTION: Therapeutic binding molecules
FILE REFERENCE: Not Yet Known
CURRENT APPLICATION NUMBER: US/11/297,317
CURRENT FILING DATE: 2005-12-08
NUMBER OF SEQ ID NOS: 30
SOFTWARE: PatentIn version 3.1
SEQ ID NO 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 9
TYPE: PRT
TYPE: PRT
CORGANISM: Artificial
PEATURE:
PEATURE:
NAME/KEY: MISC FEATURE
OTHER INFORMATION: hypervariable region
OTHER INFORMATION: ID NO:1
US-11-297-317-21
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Best Local S
Matches 4
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Pred. No. 5.2e+04;
1; Mismatches 1;
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PRIOR APPLICATION NUMBER: EP 04447239.7

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RESULT 3
US-11-122-986-778
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                                                                                    ; ORGANISM: Staphylococcus aureus
US-11-122-986-778
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US-11-140-487A-1220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: EP 05102441.2
PRIOR FILING DATE: 2005-03-25
PRIOR PELLOATION NUMBER: US 60/576,310
PRIOR PILING DATE: 2004-06-03
PRIOR PILING DATE: 2004-06-03
PRIOR APPLICATION NUMBER: US 60/622,782
PRIOR APPLICATION NUMBER: US 60/665,395
PRIOR APPLICATION NUMBER: US 60/665,395
PRIOR APPLICATION NUMBER: US 60/665,395
PRIOR APPLICATION DATE: 2005-03-25
                                                                                                                                  SEQ ID NO 778
LENGTH: 14
TYPE: PRT
                         Query Match
Best Local
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 1220
  Matches
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APPLICANT: DHARAMSI, AKIL
APPLICANT: VEDADI, MASOUD
TITLE OF INVENTION: ESSENTIAL NOVEL BACTERIAL POLYPEPTIDES
FILE REFERENCE: IPT-330.01
CURRENT APPLICATION NUMBER: US/11/122,986
CURRENT FILING DATE: 2005-05-05
PRIOR APPLICATION NUMBER: 60/423,875
PRIOR FILING DATE: 2002-11-05
PRIOR FILING DATE: 2002-11-05
                                                                                                                                                                                             Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 844
SOFTWARE: PatentIn Ver. 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/423,757
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,758
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/424,367
PRIOR FILING DATE: 2002-11-06
PRIOR FILING DATE: 2002-11-06
                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/424,373 PRIOR FILING DATE: 2002-11-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/423,832
PRIOR FILING DATE: 2002-11-05
PRIOR APPLICATION NUMBER: 60/423,915
PRIOR FILING DATE: 2002-11-05
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Local Similarity 80.0%;
                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/424,376
FILING DATE: 2002-11-06
APPLICATION NUMBER: 60/424,370
FILING DATE: 2002-11-06
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FILING DATE: 2002-11-06
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  Similarity 3; Conserv
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                      29.4%;
Score 25; DB Pred. No. 92; 3; Mismatches
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Pred. No. 5.2e+04;
                                             DB 7;
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                                         Length 14;
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    Indels
    0;
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US-10-546-594-92
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-538-066-163
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US-10-489-071-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 6, Application US/10489071
Publication No. US20060094672A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: PASQUALINE ET AL.
TITLE OF INVENTION: COMPOSITIONS AND METHODS OF USE OF TARGETING PEPTIDES AGAINST
TITLE OF INVENTION: PLACENTA AND ADIPOSE TISSUES
FILE REFERENCE: UTSC:85GUS
CURRENT FILING DATE: 2004-03-08
PRIOR APPLICATION NUMBER: PCT/US02/27836
PRIOR APPLICATION NUMBER: PCT/US02/27836
PRIOR APPLICATION UMBER: PCT/US02/27836
PRIOR FILING DATE: 2002-08-30
NUMBER OF SEQ ID NOS: 144
SOFTWARE: PATENTIN VEXISION 3.1
Sequence 92, Application US/10546594
Publication No. US20060088538A1
GENERAL INFORMATION:
APPLICANT: HOSOKKMA, Saiko
APPLICANT: AOKI, Masahiko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 767
SOFTWARE: PatentIn version 3.2
SEQ ID NO 163
LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 163, Application US/10 Publication No. US20060094649A1 GENERAL INFORMATION:
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Best Local
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TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7

TITLE OF INVENTION: Peptides and Compositions

FILE REFERENCE: 2060.015PC06

CURRENT APPLICATION NUMBER: US/10/538,066

CURRENT FILING DATE: 2005-06-09

PRIOR APPLICATION NUMBER: US 60/432,017

PRIOR FILING DATE: 2002-12-10
                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Synthetic Peptide
                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
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                                                                                                                                                                                                             3 DSPVR 7
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66.7%;
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Pred. No. 5.2e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 23; DB 6;
Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                      Mismatches
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Query Match
Best Local Similarity
Thehes 3; Conserve
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-546-594-92
RESULT 8
US-10-516-079-64
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                                                                                                                      5
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                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: EP 04012951.2
PRIOR FILLING DATE: 2004-06-01
PRIOR PELLING DATE: 2004-10-28
PRIOR APPLICATION NUMBER: EP 04447239.7
PRIOR FILLING DATE: 2004-10-28
PRIOR PELLING DATE: 2005-03-25
PRIOR PELLING DATE: 2005-03-25
PRIOR PELLING DATE: 2004-06-03
PRIOR FILLING DATE: 2004-06-03
PRIOR FILLING DATE: 2004-06-03
PRIOR APPLICATION NUMBER: US 60/622,782
PRIOR APPLICATION NUMBER: US 60/665,395
PRIOR FILLING DATE: 2005-03-25
PRIOR FILLING DATE: 2005-03-25
PRIOR FILLING DATE: 2005-03-25
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CURRENT APPLICATION NUMBER: US/10/546,594

CURRENT FILING DATE: 2005-08-19

PRIOR APPLICATION NUMBER: JP 2003/54670

PRIOR FILING DATE: 2003-02-28

PRIOR APPLICATION NUMBER: JP 2003/194643

PRIOR FILING DATE: 2003-07-09

NUMBER OF SEQ ID NOS: 132

SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1766, Application US/11140487A Publication No. US20060093617A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1 SEQ ID NO 1766
                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 2278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/11/140,487A CURRENT FILING DATE: 2005-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Innogenetics N.V.
TITLE OF INVENTION: Peptides for inducing a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: FUKUDA, KAZUMASA
TITLE OF INVENTION: MONOCLONAL ANTIBODY AND GENE ENCODING THE SAME, HYBRIDOMA, PHARMI
TITLE OF INVENTION: COMPOSITION, AND DIAGNOSTIC REAGENT
                                                                                                                                                                                                                                                                                        LENGTH: 9
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: 166
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Local Similarity 57.1%;
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KUMAI, Koichiro
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ilarity 33.3%;
Conservative
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Pred. No. 5.2e+04;
1; Mismatches 2
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Pred. No. 2.4e+02;
2; Mismatches 4
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APPLICANT: Winter, Greg
TITLE OF INVENTION: Ligand
FILE REFERENCE: 8039/2142
CURRENT APPLICATION NUMBER: US/11/023,959A
CURRENT FILING DATE: 2004-12-28
PRIOR APPLICATION NUMBER: PCT/GB03/002804
PRIOR FILING DATE: 2003-06-30
NUMBER OF SEO ID NOS: 187
SOFTWARE: Patentin version 3.3
SEQ ID NO 63
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 51530-5006-WO
CURRENT APPLICATION NUMBER: US/10/516,079
CURRENT FILING DATE: 2004-11-29
PRIOR APPLICATION NUMBER: US 60/406,033
PRIOR FILING DATE: 2002-08-27
PRIOR APPLICATION NUMBER: US 60/384,171
PRIOR FILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 95
SOFTWARE: PatentIn version 3.2
SEQ ID NO 64
            US-10-981-300-34
Sequence 34, Application US/10981300
Publication No. US20060093599A1
GENERAL INFORMATION:
APPLICANT: GIORGIO SENALDI
APPLICANT: GADI GAZIT-BORNSTEIN
APPLICANT: GADI GAZIT-BORNSTEIN
TITLE OF INVENTION: FOR MAKING AND USING THE FILE REFERENCE: ABGX-005
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US-11-023-959A-63
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Publication No. US20060106203A1
GENERAL INFORMATION:
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Publication No. US20060088899A1
GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT: GRIMES, Carol A.
APPLICANT: GONDA, Matthew A.
APPLICANT: GONDA, Combination chemotherapy with chlorotoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 10
TYPE: PRT
CURRENT APPLICATION NUMBER: US/10/981,300
                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
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Pred. No. 3.2e+02;
1; Mismatches 1
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Pred. No. 5
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                                                        AND METHODS
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Sequence 84, Application US/11106014
Publication No. US20060088846A1
GENERAL INFORMATION:
APPLICANT: Pagano, Michele
APPLICANT: Chiaux, Dah Sharim
APPLICANT: Chiaux, Dah Sharim
APPLICANT: Latres, Esther
APPLICANT: Szivastava, Promod
APPLICANT: Chandawarker, Rajiv
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US-11-140-487A-2207
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; TYPE: PRT
; ORGANISM: homo sapiens
US-10-981-300-34
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Sequence 2207, Application US/11140487A

Publication No. US20060093617A1

GENERAL INFORMATION:

APPLICANT: Innogenetics N.V.

APPLICANT: Innogenetics of inducing a CTL and/or HTL response to Hepatitis C virithe Reference: 166

CURRENT APPLICATION NUMBER: US/11/140,487A

CURRENT FILING DATE: 2005-05-31

PRIOR APPLICATION NUMBER: EP 04012951.2
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NUMBER OF SEQ ID NOS: 71
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 34
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CURRENT FILING DATE: 2005-04-13
PRIOR APPLICATION NUMBER: 10/632,150
PRIOR FILING DATE: 2003-07-30
PRIOR APPLICATION NUMBER: 10/042,417
PRIOR FILING DATE: 2002-01-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/124,449 PRIOR FILING DATE: 1997-03-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 09/385,219
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: 60/118,568
PRIOR FILING DATE: 1999-02-03
PRIOR APPLICATION NUMBER: 60/098,355
PRIOR FILING DATE: 1998-08-28
PRIOR FILING DATE: 1998-08-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Chandawarker Rajiv
TITLE OF INVENTION: METHODS TO IDENTIFY COMPOUNDS USEFUL FOR THE TREATMENT
TITLE OF INVENTION: OF PROLIFERATIVE AND DIFFERENTIATIVE DISORDERS
FILE REFERENCE: 5914-106-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 12
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
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Similarity 44.4%;
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Pred. No. 5.8e+02;
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Pred. No. 5.3e+02;
2; Mismatches 4
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US-10-538-066-159
; Sequence 159, Application US/10538066
; Publication No. US20060094649A1
; GENERAL INFORMATION:
; APPLICANT: Epimmune Inc.
; TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7, &
; TITLE OF INVENTION: Peptides and Compositions
; FILE REFERENCE: 2060.015PC06
; CURRENT APPLICATION NUMBER: US/10/538,066
; CURRENT APPLICATION NUMBER: US 60/432,017
; PRIOR APPLICATION NUMBER: US 60/432,017
; PRIOR FILING DATE: 2002-12-10
; NUMBER OF SEQ ID NOS: 767
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 767
SOFTWARE: PatentIn version 3.2
SEQ ID NO 404
LENGTH: 8
                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.1 SEQ ID NO 2207
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Epimmune Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7
TITLE OF INVENTION: Peptides and Compositions
FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 2004-10-29
PRIOR APPLICATION NUMBER: US 60/665,395
PRIOR FILING DATE: 2005-03-25
NUMBER OF SEQ ID NOS: 2278
NUMBER OF SEQ ID NOS: 2278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/576,310 PRIOR FILING DATE: 2004-06-03 PRIOR APPLICATION NUMBER: US 60/622,782 PRIOR FILING DATE: 2004-10-29
SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: EP 05102441.2
PRIOR FILING DATE: 2005-03-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: EP 04447239.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
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o. US20060094649A1
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Pred. No.
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Sequence 429, Application US/10538066

Publication No. US20060094649A1

GENERAL INFORMATION:
APPLICANT: Epimmune Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7, and -B44 Tumor Associated Antigen
TITLE OF INVENTION: Deptides and Compositions
FILE REFERENCE: 2060.015PC06

CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR APPLICATION NUMBER: US 60/432,017
SOFTWARE: PatentIn version 3.2
SEQ ID NO 429
LENGTH: 9
LENGTH: 9
LENGTH: Homo sapiens
US-10-538-066-429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; SEQ ID NO 159
; LENGTH: 9
; TYPE; PRT
; ORGANISM: Homo sapiens
US-10-538-066-159
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Search completed: May 30, 2006, 10:25:03 
Job time : 5.66667 secs
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US-10-538-066-429
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                                                                                                                                                           Query Match 22.4%; Score 19; DB 6; Length 9; Best Local Similarity 57.1%; Pred. No. 5.2e+04; Matches 4; Conservative 1; Mismatches 2; Indels
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length: 0
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| Abg98753  | Aec97896  | Adw23329  | Adw23210  | Adw23315  | Adv51742 | Adv51741 | Adv51740 | Adv51738 | Adv51743 | Adv51739 | Adv51733 | Adv51736 | Adv51734 | Adv51737 | Adv51735 | Aae05129  | Aea45250 | Adz86156  | Aby01210  | Aby01523  | Abj19566 |
| F protein | HLA-A24-b | SARS coro | SARS coro | SARS coro | SARS-CoV | Peptide # | dodi     | SARS coro | SARS coro | SARS coro | Neuronal |

ALIGNMENTS

#### ADR10601 RESULT 1 Dog IgE epitope recognised by monoclonal antibody 5.91, SEQ ID 1. Canis familiaris. Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody. ADR10601; 16-JAN-2003; 2003US-0440472P. 15-JAN-2004; 2004WO-US003566 05-AUG-2004. WO2004065936-A2 21-OCT-2004 ADR10601 standard; peptide; 15 (UYNC-) UNIV NORTH CAROLINA STATE (first entry) Ş

Hammerberg B;

Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian I or treating asthma or anaphylactic shock. IgE

WPI; 2004-593545/57.

Example 6; Page 9; 14pp; English.

The present invention relates to a novel monoclonal antibody (I) that specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. (I) is useful for testing an allergen reactivity of an IgE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut and corn allergens. The sample is a biological sample collected from a dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating sathma or anaphylactic shock. The 5-91 and 3.76 monoclonal antibodies recognise epitopes on canine IgE corresponding to amino acid residues 357-371 (ADR10601) and 146-162 (ADR10603) respectively of the IgE epsilon-chain. Recognition of epsilon-chains for

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RESULT 2
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                                                                                                                                          CC The present invention relates to a novel monoclonal antibody (I) that CC specifically binds to a mammalian IgE epitope, where the epitope is CC between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. CC (I) is useful for testing an allergen reactivity of an IgE sample. The CC allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut CC and corn allergens. The sample is a biological sample collected from a CC dog, cat or horse. (I) is also useful for detecting mammalian IgE and for CC treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal CC antibodies recognise epitopes on canine IgE corresponding to amino acid CC residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the CC canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from CC at, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and CC 3.76 were observed to have good cross-reactivity with the epsilon-chain CC of IgE from cat and horse, but did not exhibit cross-reactivity with CC either pig or human epsilon-chains of IgE. The present sequence is the CC pig IgE 5.91 recognition site.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian I or treating asthma or anaphylactic shock.
                                                                                            Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody;
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l Similarity 100.0%;
15; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         UNIV NORTH CAROLINA STATE
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65.4%;
57.1%;
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   Score
Pred.
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Pred. No. 5.9e-07;
Mismatches 0;
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   No. 53;
   DB 8;
0.044;
                                 Length 15;
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4

RESULT 4 ADR10603

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RESULT 3
ADR10602
ID ADR1
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                                                                                                                                                                                                                                                                                                 The present invention relates to a novel monoclonal antibody (I) that CC specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. CC (I) is useful for testing an allergen reactivity of an IgE sample. The callergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut CC allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut CC and corn allergens. The sample is a biological sample collected from a CC dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal CC treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal CC catious seepilon-chain. Recognition of Egg corresponding to amino acid CC catious 357-371 (ApR10601) and 146-162 (ApR10609) respectively of the CC canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from CC cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and CC cat, horse, pig and horse, but did not exhibit cross-reactivity with the epsilon-chain epsilon-chains of IgE from cat and horse, but did not exhibit cross-reactivity with the epsilon or human epsilon-chains of IgE. The present sequence is the
                                                                                                        Query Match
Best Local S
Matches 8
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                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian IgE or treating asthma or anaphylactic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 6; Page 9; 14pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cat IgE epitope recognised by monoclonal antibody 5.91,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-OCT-2004
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                                                  N
                                                                                                           8; Conserv
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NDSPVRTEQQATT 14
                                                  NDSPIQTDQYTTT 14
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                                                                                                              Conservative
                                                                                                                                                                                                                                                                                  recognition site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry
                                                                                                                                         61.5%;
                                                                                                                                                                   59.3%;
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                                                                                                                                         Score 48; D
Pred. No. 0.
                                                                                                                 Mismatches
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                                                                                                                                         DB 8;
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7.
                                                                                                                                                                Length 15;
                                                                                                                 Indels
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RESULT 5
ADC64568
ID ADC6
XX
AC ADC6
XX
AC ADC6
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AC HOT6
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DT 18-E
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DE HOT6
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HOT6
                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Thes 9; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     specifically binds to a mammalian IgE epitope, where the epitope is between amino acids 145-166 or 356-374 of mammalian IgE, e.g. dog IgE. (I) is useful for testing an allergen reactivity of an IgE sample. The allergen is chosen from mold, pollen, dust mite, milk, egg, soy, peanut and corn allergens. The sample is a biological sample collected from a dog, cat or horse. (I) is also useful for detecting mammalian IgE and for treating asthma or anaphylactic shock. The 5.91 and 3.76 monoclonal antibodies recognise epitopes on canine IgE corresponding to amino acid residues 357-371 (ADR10601) and 146-162 (ADR10609) respectively of the canine IgE epsilon-chain. Recognition of epsilon-chains from IgE from cat, horse, pig and human by 5.91 and 3.76 was examined. Both 5.91 and and in the case of the canine IgE and human by 5.91 and 3.76 was examined.
                                                                                                                                                                                                                                                                                                                                                                                                                 3.76 were observed to have good cross-reactivity with the epsilon-chain of IgE from cat and horse, but did not exhibit cross-reactivity with either pig or human epsilon-chains of IgE. The present sequence is the horse IgE 5.91 recognition site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel antibody that specifically binds to mammalian IgE epitope, useful for testing an allergen reactivity of IgE sample, detecting mammalian IgE or treating asthma or anaphylactic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antiasthmatic; Antiallergic; Immunosuppressive; IgE; dog; asthma; anaphylactic shock; 5.91 monoclonal antibody; 3.76 monoclonal antibody;
                                   Horse immunoglobulin E, IgE, heavy chain immunogenic peptide P4
                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 6; Page 9; 14pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hammerberg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Horse IgE epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-OCT-2004
Horse; immunoglobulin E; IgE; heavy chain; immunogen; allergy
                                                                       18-DEC-2003
                                                                                                                                             ADC64568 standard; peptide; 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADR10603 standard; peptide; 15 AA
                                                                                                           ADC64568
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             present invention relates to a novel monoclonal antibody (I) that
                                                                                                                                                                                                                                                                        RNDSPIQTDQYTTT 14
                                                                                                                                                                                                                                      RNNVLIOTDOOATT 14
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Ş
                                                                                                                                                                                                                                                                                                           Conservative
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                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         recognised by monoclonal antibody 5.91,
                                                                                                                                                                                                                                                                                                                            50.6%;
                                                                                                                                                                                                                                                                                                         Score 41; DB 8; Length 15;
Pred. No. 5.4;
1; Mismatches 4; Indels
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                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                           Gaps
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CC isolated polypoptide having an amino acid sequence that is at least 80% cc identical to 6 (S1-S6), 15 amino acid sequence that is at least 80% cc identical to 6 (S1-S6), 15 amino acid sequences derived from cc quine immunoglobulin [G]E), the six composition induces production of an expectation of the six cc polypoptides are not explicitly identified in the specification. Also cc included are a composition comprising an antibody that specifically binds to equine immunoglobulin (Ig)E), the six composition are polypoptide at least 80% identical to (S1)-(S6), an antibody that specifically binds to equine IgE made by the process of immunising an animal with a composition further comprising an isolated polypoptide (the amino acid sequence of the polypoptide is at least 80% identical to (S1)-(S6), making an cc amino acid sequence of the polypoptide is at least 80% identical to (S1)-(S6), and collecting antiserum from the animal) and a kit for detection cof equine IgE in a biological sample comprising the antibody and means cf of detecting specific binding of the antibody to equine IgE. The cample (serum) which involves contacting the sample with the antibody, the sample with the antibody. Camplex can be detected. The complex. The antibody and antigen are complex can be detected. The complex is detected using a second labelled cantibody. The poptides are useful for generating antibodies specific for IgE which can serve as a diagnostic test for allergy. The present cc poptide from the early portion of the C4 region.
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunogenic composition comprising an isolated equine immunoglobulin E polypeptide that induces production of antibodies which specifically bind to equine immunoglobulin E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-NOV-2001; 2001US-00052788
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 8; 14pp; English.
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                                                              Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to an immunogenic composition comprising an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (REGC ) UNIV CALIFORNIA
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  42.0%;
77.8%;
  Score 34; DB Pred. No. 89;
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                       7;
                       Length 15;
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                6 IQTDQYTTT 14
IQTDQQATT 9
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맑 S

Matches

Conservative

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Indels

0

Gaps

0

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RESULT 6
ABG98754
                           ABG98754;
13-JAN-2003
                                                    ABG98754 standard; peptide; 10 AA
(first entry)
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F protein; RSV; decapeptide; negative sense single stranded RNA virus; viral F protein; F protein mediated host cell virion fusion; virucide;

F protein decapeptide #241.

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RESULT 7
ABG98755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC protein or its derivative from a negative sense single stranded RNA CC virus, by expressing a nucleic acid molecule encoding the protein in a CC host cell, where the nucleic acid is optimised for expression by a CC eukaryotic cell. The protein, especially F protein, is useful for CC expression, designing and/or modifying an agent capable of interacting CC with a viral F protein or its derivative and modulating a functional CC activity associated with the protein, by contacting the protein with a CC putative agent and assessing the degree of interactive complementarity of CC the agent with the protein. An optimised nucleic acid or its derivative, CC equivalent, analogue or mimetic is useful for interacting with a viral F CC protein and modulating a functional activity associated with the viral CC protein, for use in the manufacture of a medicament utilised in the CC therapeutic and/or prophylactic treatment of conditions characterised by infection with a negative sense single stranded RNA virus, and for CC modulating a functional activity associated with a viral F protein in a CC subject, preferably a mammal, especially a human, where the functional CC subject, preferably a mammal, especially virion fusion and/or virion budding and the modulating is down regulation. Sequences ABG98514-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
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                                                        F protein; RSV; decapeptide; negative sense single stranded RNA virus; viral F protein; F protein mediated host cell virion fusion; virucide; F protein mediated host cell virion budding.
Respiratory syncytial virus
                                                                                                                                                                                          F protein
                                                                                                                                                                                                                                                             13-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                  ABG98755 standard; peptide; 10 AA
                                                                                                                                                                                                                                                                                                                         ABG98755;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-NOV-2000; 2000US-0252767P
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                                                                                                                                                                                          decapeptide #242.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nod for facilitating production of a a negative sense single stranded RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .3e+02;
3;
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RESULT 8
AAR79898
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 6
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                                                                                                                                          Respiratory syncitial virus; RS virus; fusion protein; antibody; vaccine;
                                                                                                                                                                        Fusion protein synthetic fragment SP4.
                                                                                                                                                                                                             25-MAR-2003
25-APR-1996
                                                                                                                                                                                                                                                                                                    AAR79898 standard; peptide; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 42; Page 102; 367pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    a eukaryotic cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-NOV-2001; 2001WO-AU001517
                                                                                                                        human; therapy; diagnosis; immunoassay
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                6;
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                                                                                                                                                                                                             (revised)
(first entry)
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Pred. No. 1.3e+02;
Pred. No. 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 10;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antibodies against these fragments could be used to confer short-term protection against the RS virus. The vaccine can then be used to protect humans and animals (for animals, the RS sequence used is from the RS virus of that species). The dose of the vaccine is 0.1-100 micro grammes per kg body weight. The vaccine can also be used in diagnosis, such as immunoassays. The DNA encoding this sequence can be used in hybridisation tests to detect RS virus in blood. The antibodies can be used to identify viral epitopes. The advancage with using this methodology is that there is no enhancement/potentiation of the disease seen with current treatments. (Updated on 25-MAR-2003 to correct PF field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Paradiso
Walsh EE;
                                                                                                                                                                                Rat; glial cell line-derived neurotrophic factor; GDNF; receptor; treatment; dopaminergic nerve cell disorder; Parkinson's disease; Alzheimer's disease; amyotrophic lateral sclerosis; diabetes; Huntington's disease; glaucoma; retinal degeneration; hearing loss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The sequences represented by AAR79895-R79899 are synthetic peptides corresponding to regions of the respiratory syncitial (RS) virus fusion protein (see AAR79894). This sequence corresponds to residues 261-273. These fragments were used to create antibodies. The DNA encoding these fragments is inserted into expression vectors and used to transform cell cultures, the expressed protein is then isolated and purified. These peptide fragments are then used in a vaccine. Alternatively, the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sub-unit vaccine against respiratory syncytial polypeptide related to neutralising epitope on
                                                                                                                                        Rattus
                                                                                                                                                                     gene therapy
                                                                                                                                                                                                                                                               Rat GDNF receptor
                                                                                                                                                                                                                                                                                             01-MAY-1998
                                                                                                                                                                                                                                                                                                                                                       AAW35339 standard; peptide; 14 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13
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20-SEP-1989;
 22-APR-1996;
09-MAY-1996;
                                                                            30-OCT-1997.
                                                                                                         WO9740152-A1
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                                              15-APR-1997;
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89US-00409915.
96US-0015907P
96US-0017221P
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                                                                                                                                                                                                                                                                residues
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Pred. No. 1.7e+02;
0; Mismatches 3
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G protein,
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pharmaceutical formulations. Receptor expressing cells, preferably transfected ex vivo, can be used similarly by implantation, and the use of the receptor cDNA in gene therapy is also contemplated. Probes based on the cDNA can be used to identify GDNF responsive cells and tissues, e.g. to identify patients who would benefit from GDNF therapy, and abnormalities in receptor expression, and to isolate molecules that form a complex with the cDNA or are homologous/cross-reactive with the cDNA. Anti-receptor antibodies, oligonucleotides derived from the cDNA and animal models that overexpress the receptor can be used to study the biological function of GDNF, knockout transgenic animals can be used to detect GDNF dependent neurons or processes and the antibody can be used in immunoassays for the receptor. The receptor binds GDNF specifically and with high affinity, acting as part of a complex that mediates/enhances signal transduction by GDNF, i.e. increasing dopamine
                                                                                                                                                                                                                                                                                                                                                                                                                                     Huntington's disease and (optionally in combination with GDNF) glaucoma, retinal degeneration and hearing loss caused by injury to inner ear sensory neurons. The receptor can also be used to block unwanted GDNF activity, analyse GDNF related molecules and stabilise GDNF in pharmaceutical formulations. Receptor expressing cells, preferably transfected ex vivo, can be used similarly by implantation, and the use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present peptide is derived from rat glial cell line-derived neurotrophic factor (GDNF) receptor, which can be used to treat dopaminergic nerve cell disorders, e.g. Parkinson's and Alzheimer's disease or amyotrophic lateral sclerosis, complications of diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Glial cell line derived dopaminergic nerve cell
                                                                                           Sequence 14
                                                                                                                                        uptake in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-APR-1997;
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                                                                                                                                      dopaminergic cells
                                                                                               AA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 21; 196pp; English
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                       38.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neurotrophic factor receptor - useful to treat disorders, e.g. Parkinson's and Alzheimer's
Score 31; DB 2; Le
Pred. No. 2.8e+02;
w:ematches 3;
                                           Length 14;
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                                Similarity
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PVOTTTATTT
                          Conservative
13
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25-MAR-1999
                                                                           AAW84178 standard; peptide; 14
(first entry)
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03-DEC-1998 WO9854213-A2 Synthetic.

gp.

retinal ganglion

cell

degeneration;

sensory

neuropathy;

retinopathy;

therapy

Rat; glial cell-line derived neurotrophic factor receptor-alpha; glial cell line-derived neurotrophic factor; GDNF; neurturin; signal transduction; dopaminergic nerve cell; Parkinson's disease; Alzheimer's disease; amyotrophic lateral sclerosis; neurological disorder; diabetes; glaucoma; sensory neuron;

Rat GDNFR-alpha derived peptide.

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RESULT 11

AEC98667
ID AEC98
XX AEC98
XX O1-DE
XX O1-DE
XX HLMA-[
XX Humar
XX epitc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                31-DEC-2003;
02-JUL-2004;
                                                                                                             03-JAN-2005; 2005WO-US000077
                                                                                                                                                                                                                                                                                     Human papillomavirus.
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                                                                                                                                                                                                                                WO2005089164-A2
                                                                                                                                                                                                                                                                                                                                                                     epitope
                                                                                                                                                                                                                                                                                                                                                                                                  Human papilloma virus infection; virucide; cancer; cytostatic; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       HLA-DR binding
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                                                                                                                                                                                                                                                                                                                                          leukocyte
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                     mapping; immune stimulation; cytotoxic T-lymphocyte;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             scierosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                2003US-0533211P
2004US-0584652P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                       epitope
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                                                                                                                                                                                                                                                                                                                                             antigen.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 31; DB 2;
Pred. No. 2.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                             #156.
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The invention relates to a new polynucleotide comprising a multi-epitope construct comprising nucleic acids encoding the human papillomavirus (HPV) cytotoxic T lymphocyte (CTL) epitopes e.g., HPV16.El. 214, and that care directly or indirectly joined to one another in the same reading cframe, a vaccine minigene. Also included are a vector comprising the multi-epitope construct, a polypeptide comprising a maino acid sequence encoded by the polynucleotide, a composition (comprising the polynucleotide, vector and/or polypeptide and a carrier), a cell comprising the polynucleotide, vector or polypeptide, inducing an immune response against human papillomavirus virus (HPV) and making the colynucleotide, vector or polypeptide. The epitopes are derived from the Ell E2, E6 and E7 proteins. The epitopes may be linked via a GP-anchor/spacer peptide. The order of the epitopes may be linked via a GP-anchor/spacer peptide. The order of the epitopes in the vaccine protein are disposed in the tables referred to in the claims of the specification. The polynucleotide, vector or polypeptide is useful in preparing a composition for inducing an immune cresponse against human papillomavirus virus (HPV) and thus providing a defense against human papillomavirus virus (HPV) and thus providing a captore is an HPV-derived peptide epitope that binds to a subclass of human leukocyte antigen (HLA) peptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chesnut Chen Y,
Sequence
                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 333;
                                                                                                                                                                                                                                                                                                                                                                                                                              New polynucleotide comprises a multi-epitope construct comprising nucleic acids encoding the human papillomavirus (HPV) cytotoxic T lymphocyte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2005-658982/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (BABE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BAKE/)
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                                                                                                                                                                                                                                                                                                                                                                                                              epitopes, useful in preparing a vaccine against HPV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      . Deyoung
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BABE L
CHEN Y.
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BAKER D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NEWMAN M J.
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BABE L M.
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INNOGENETICS NV.
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                                                                                                                                                                                                                                                                                                                                                                                 518pp; English
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Power
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            4
                          Similarity 55. 5; Conservative
            SPIQTDQYT
SPYKIDEWI
10
            12
                                38.3%;
55.6%;
                                Score 31;
Pred. No.
                          Mismatches
                                 DB 9;
3e+02;
                          ۲.
                                      Length 15;
                          Indels
                          0
                          Gaps
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AAM98177 standard; peptide; 14

24-JAN-2002 (first entry)

Human peptide #1452 encoded by a SNP oligonucleotide.

RESULT 12
AAM98177
ID AAM98
XX AAM98
AC AAM98
XX 24-JA
DT 24-JA
DT 24-JA
DT 1munn
XX Immun
XX Immun
XX Immun
XX amyloo
KW neuro
KW annyloo
KW compl
KW compl
KW intex
KW intex
KW intex neuroprotective; antimicrobial; gene therapy; vaccine; amylase; can amyloid protein; angiopoletin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating tomplement related protein; cytochrome; kinase; cytokine; interferinterfleukin; G-protein coupled receptor; thioesterase; inflammation Immunosuppressive; immunostimulatory; antiinflammatory; infection; inflammation; cytostatic; cancer;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC The present invention relates to oligonucleotides (see AAL26793-AAL34659) CC encoding polymorphic variants of proteins related to amylases, amyloid CC proteins, angiopoietin, apoptosis related proteins, cadherin, cyclin, CC polymerase, oncogenes, histones, kinases, colony stimulating factors, CC interferons, interleukins, G-protein coupled receptors and thioesterases. CC The present sequence is a peptide encoded by one such oligonucleotide. CC The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with CC inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases (c) arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, branch, breast, colon and kidney, leukaemia), diseases of the nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-DEC-1999; 99US-0173419P
27-DEC-2000; 2000US-00173419
                                                                                                                      Neuroprotective; Vulnerary; Vasotropic; Cerebroprotective; spinal cord injury; brain injury; paralysis; neurodegenera
                                                                                                                                                                                                                                                                  AED12588 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             brain, breast, colon and kidney, leukaemia), dissystem and an infection of pathogenic organisms
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-465210/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200147944-A2
                                                                                                        cerebrovascular
                                                                                                                                                                      Anti-NogoR scFv protein NGF0121 HC CDR3 sequence
                                                                                                                                                                                                      01-DEC-2005
                                                                                                                                                                                                                                     AED12588;
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              29-SEP-2005
                                            US2005215770-A1
                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                   VQTDKVTST 13
                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                      (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leach M;
                                                                                                       ischemia; Nogo receptor; NogoR.
                                                                                                                                                                                                                                                                                                                                                                                                                              37.0%;
                                                                                                                       injury; paralysis; neurodegenerative disease;
                                                                                                                                                                                                                                                                     14
                                                                                                                                                                                                                                                                                                                                                                                                               Score 30; DB 4; Length 14;
Pred. No. 4.1e+02;
3; Mismatches 1; Indels
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RESULT 14
AAY51451
ID AAY51
XX AAY51
XX AAY51
XX T1-MA
XX AAV V
XX VP3;
XX VP3;
XX VP3;
XX VP3;
XX VP3;
XX I9-DE
XX Adeno
XX MC996
XX I9-DE
XX I9-JU
XX I9-JU
XX IP-JU
XX IP-J
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a novel antibody or its fragment (e.g. an scFv) that binds to the Nogo receptor (NogoR; AED12284). The antibody comprises a Variable Heavy Complementarity Determining Region (VHCDR). VHCDR2 or VHCDR3 and a Variable Light Complementarity Determining Region (VLCDR). VLCDR2 or VLCDR3. The antibody prevents binding or inhibits interaction of NogoR with: p75(NTR) (AED12288); INGO-1 (AED12289); Nogo (AED12285); OMGP (AED12286); or MAG (AED12287), and promotes neurite outgrowth or axonal regeneration. The antibody is useful in preparing a composition for treating or ameliorating spinal cord injury, brain trauma, paralysis, neurodegenerative disorders or stroke. The present sequence is a CDR peptide sequence for one such scFv that binds to NogoR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antibody or its fragment that specifically binds Nogo receptor (NogoR), useful in preparing a composition for treating or ameliorating spinal cord injury, brain trauma, paralysis, neurodegenerative disorders or stroke.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14 AA;
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                               Example 2; Page 42; 43pp; German
                                                                                            Structure proteins of adeno-associated virus with at least one mutation have increased infectiveness.
                                                                                                                                                                                     WPI; 2000-160586/14.
                                                                                                                                                                                                                                           Hallek M, Ried M,
                                                                                                                                                                                                                                                                                                                                                                    19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Adeno-associated virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VP3; structural protein; capsid; tropism; diagnosis; gene therapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAY51451;
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                                                                                                                                                                                                                                              Deleage G,
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45.5%;
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Pred. No. 4.1e+02;
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CC This invention describes a novel structural protein (I) of adeno-
CC associated virus (AAV) which has at least one mutation that alters the
CC chromatographic properties of the virus. The invention also describes (1)
CC anucleic acid (II) that encodes (I); (2) a cell (III) containing (II);
CC and (3) producing (I) comprising culturing (III). (II) and (III) are
CC used to purify AAV and its particles, to alter the tropism or
CC antigenicity of AAV, to transform cells, for genomic targeting, for
CC diagnosis, for examination of activity and/or gene therapy. Altering the
CC concentration of virus and produces viral particles to a higher titre or
higher purity and/or provides more efficient purification, especially
CC and a quicker and less expensive process). Particularly, mutated (I)
CC concentration than the wild type, which tends to co-elute with other
CC viruses, serum proteins and cellular debris, or that include a specific
CC affinity tag. Mutation may also allow targeting of cells, e.g.
CC hematopoietic cells, that are not normally targets for AAV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VP3 protein; structural protein; chromatography; tropism; antigenicity;
genomic targeting; gene therapy; purification; affinity tag.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 15; 20pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New mutant adeno-associated virus structural protein with altered chromatographic properties, useful in preparation of gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-203661/21.
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| KMC_Celerra_SIDSJ/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
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| KMC_Celerra_SIDSJ/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*
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                                              Sequence 7, Application US/10758165
; Publication No. US20050196816A1
; GENERAL INFORMATION:
APPLICANT: Hammerberg, Bruce
; TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
; FILE REFERENCE: 5051-661
; CURRENT APPLICATION NUMBER: US/10/758,165
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: US 60/440,472
; PRIOR FILING DATE: 2003-01-16
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 7
; LENGTH: 15
; TYPE: PRT
; ORGANISM: Sus scrofa
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; TYPE: PRT
; ORGANISM: Canis familiaris
US-10-758-165-1
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US-10-758-165-7
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Publication No. US20050196816A1
GENERAL INFORMATION:
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Best Local Similarity 100.0%;
Matches 15; Conservative 0
   Query Match
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     65.4%;
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|  |  | US-10-530-061-2185 |      |      |      |      | US-10-279-991-13 | US-10-078-968-5 | US-10-475-049A-15 | US-10-432-234A-251 | US-10-149-138-1902 | US-10-149-138-1265 |          | US-10-149-138-1902 | US-10-149-138-1265 | US-10-149-138-543 |      |      | CC FC 0.0 |

ALIGNMENTS

### APPLICANT: Hammerberg, Bruce TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN FILE REFERENCE: 5051-661 CURRENT APPLICATION NUMBER: US/10/758,165 CURRENT FILING DATE: 2004-01-16 PRIOR APPLICATION NUMBER: US 60/440,472 PRIOR FILING DATE: 2003-01-16 NUMBER OF SEQ ID NOS: 16 SOFTWARE: PatentIn version 3.2 SEQ ID NO 1 <u>,</u> Score 81; DB 5; Pred. No. 5.4e-07; ); Mismatches 0; Ä Length 15 MAMMALIAN SPECIES 0 Gaps

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Score 53;

DB <u>ა</u>

Length 15;

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APPLICANT: Hammerberg, Bruce
ITITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR FILING DATE: 2003-01-16
NUMBER: US 60/440,472
PRIOR FILING DATE: 2003-01-16
NUMBER: OF SEQ ID NOS: 16
SOFTWARE: Patentin version 3.2
SEQ ID NO 2
LENGTH: 15
TYPE: PRT
ORGANISM: Felis catus
US-10-758-165-2
                                                                                                                                                                                                                                                                                                                                                    Sequence 3, Application US/10758165

Publication No. US20050196816A1

GENERAL INFORMATION:
APPLICANT: Hammerberg, Bruce
TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN
FILE REFERENCE: 5051-661
CURRENT APPLICATION NUMBER: US/10/758,165
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR FILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn version 3.2
SEQ ID NO 3
                                                                               RESULT 5
US-10-052-788-4
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; TYPE: PRT
; ORGANISM: Equus caballus
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Best Local Similarity
Ahes 8; Conserv
                              Sequence 4, Application US/10052788 Publication No. US20030087314A1 GENERAL INFORMATION:
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Publication No. US20050196816A1
GENERAL INFORMATION:
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APPLICANT: Gershwin, Laurel J. APPLICANT: Pettigrew, Howard David
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Sequence 250, Application US/10432234A
Publication No. US20040161846A1
GENERAL INFORMATION:
APPLICANT: Biota Scientific Management Pty LtyTITLE OF INVENTION: A method of expression and FILE REFERENCE: 12084720/TDO
CURRENT APPLICATION NUMBER: US/10/432,234A
CURRENT FILING DATE: 2003-05-22
PRIOR APPLICATION NUMBER: US 60/252767
PRIOR FILING DATE: 2000-11-22
NUMBER OF SEQ ID NOS: 574
SOFTWARE: Patentin version 3.1
SEQ ID NO 250
LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Biota Scientific Management Pty Ltd
TITLE OF INVENTION: A method of expression and age
FILE REFERENCE: 12084720/TDO
CURRENT APPLICATION NUMBER: US/10/432,234A
CURRENT FILING DATE: 2003-05-22
PRIOR APPLICATION NUMBER: US 60/252767
PRIOR FILING DATE: 2000-11-22
NUMBER OF SEQ ID NOS: 574
SOFTWARE: PatentIn version 3.1
SEQ ID NO 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; LENGTH: 10
; TYPE: PRT
; ORGANISM: respiratory syncytial virus
US-10-432-234A-249
                                                                                                                                                                                                                                                                                                            US-10-432-234A-250
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US-10-432-234A-249
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Kalina, Warren V.
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Epsilon Immunoglobulin Chain Derived Peptides
TITLE OF INVENTION: Induction of Anti-IgE Antibodies
FILE REFERENCE: 023070-121000US
FILE REFERENCE: 023070-121000US
CURRENT FAPPLICATION NUMBER: US/10/052,788
CURRENT FILING DATE: 2001-11-08
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence:epitope peptide OTHER INFORMATION: P4, early portion of C4 of equine IgE epsilon OTHER INFORMATION: heavy chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 15
TYPE: PRT
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6; Conserva
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milarity 66.7%;
Conservative (
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Pred. No. 1.1e+02;
Pred. No. 3;
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Pred. No. 76;
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; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 29
; LENGTH: 14
; TYPE: PRT
; ORGANISM: RAT
US-10-155-693-29
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                                                            CURRENT APPLICATION NUMBER: US/10/872,161
CURRENT FILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: US/08/866,354
PRIOR FILING DATE: 1997-05-30
PRIOR APPLICATION NUMBER: US 60/015,907
PRIOR ETLING DATE: 1996-04-22
PRIOR APPLICATION NUMBER: US 60/017,221
PRIOR FILING DATE: 1996-04-14
NUMBER OF SEQ ID NOS: 61
SCOPTWARE: DATE: 1997-04-14
NUMBER: DATE: DATE: 1997-04-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 29, Application US/10872161 Publication No. US20040235714A1
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SEQ ID NO 29
LENGTH: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: GLIAL CELL LINE-DERIVED NEUROTROPHIC FACTOR RECEPTOR FILE REFERENCE: A-401C CURRENT SPELICATION NUMBER: US/10/155,693 CURRENT FILING DATE: 2002-05-24 PRIOR APPLICATION NUMBER: US/08/837,199 PRIOR FILING DATE: 1997-04-14 PRIOR PILING DATE: 1997-04-14 PRIOR PILING DATE: 1996-04-12 PRIOR APPLICATION NUMBER: US 60/015,907 PRIOR APPLICATION NUMBER: US 60/015,907 PRIOR PILING DATE: 1996-04-22 PRIOR PILING DATE: 1996-05-09 PRIOR FILING DATE: 1996-05-09 NUMBER: US 60/017,221 PRIOR PILING DATE: 1996-05-09 NUMBER: US 60/017,221 PRIOR PILING DATE: 1996-05-09
                                                                                                                                                                                                                                                                                                                                    APPLICANT: FOX, GARY M.
APPLICANT: JING, SHUQIAN
APPLICANT: WEN, DUANZHI
TITLE OF INVENTION: GLIAL CELL LINE-DERIVED NEUROTROPHIC FACTOR RECEPTOR
FILE REFERENCE: A-401D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: FOX, GARY M.
APPLICANT: JING, SHUQIAN
APPLICANT: WEN, DUANZHI
                                                  SOFTWARE: PatentIn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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les 6; Conser
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1; Mismatches
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; ORGANISM: RAT US-10-872-161-29
                                                                              ; ORGANISM: Artificial Sequence ; FEATURE: OTHER INFORMATION: Synthetic construct US-10-601-953-561
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US-10-462-452-433
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US-10-462-452-433
                                                                                                                                                             APPLICATION., Steven C.

APPLICANT: Quay, Steven C.

APPLICANT: Quay, Steven C.

TITLE OF INVENTION: Compositions And Methods For Modulating Physiology Of Epithelial TITLE OF INVENTION: Unctional Adhesion Molecules For Enhanced Mucosal Delivery Of TITLE OF INVENTION: Therapeutic Compounds

FILE REFERENCE: 02-0319

CURRENT APPLICATION NUMBER: US/10/601,953

CURRENT APPLICATION NUMBER: 60/392,512

PRIOR APPLICATION NUMBER: 60/392,512

PRIOR APPLICATION NUMBER: 60/392,512

PRIOR APPLICATION NUMBER: 60/392,512

SOFTWARE: PatentIn version 3.2

SOFTWARE: PatentIn version 3.2

SEQ ID NO 561

LENGTH: 10
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Publication No. US20040077540A1
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  Matches
                                        Query Match
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Best Local
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CURRENT FILING DATE: 2003-06-16
PRIOR APPLICATION NUMBER: 60/393,066
PRIOR FILING DATE: 2002-06-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Compositions and Methods for Enhanced TITLE OF INVENTION: Mucosal-Delivery of Interferon Beta FILE REFERENCE: 02-02US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Quay Steven
APPLICANT: El Shafy, Mohammed Abd
APPLICANT: Gupta, Malini
APPLICANT: de Meireles, Jorge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
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Local Similarity
les 5; Conserv
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). US20040037809A1
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  Conservative
                     35.8%;
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                     Score 29; DB 4;
Pred. No. 3.6e+02;
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Pred. No. 3.6e+02;
1; Mismatches 2;
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Pred. No. 2.3e+02;
  Mismatches
                                        Length 10;
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  Indels
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APPLICANT: University Court of the University of Glasgow TITLE OF INVENTION: Targeting peptides FILE REFERENCE: PC/MC/JM/p1910US CURRENT APPLICATION NUMBER: US/09/990,832C CURRENT FILING DATE: 2003-01-27 NUMBER OF SEQ ID NOS: 127 SOFTWARE: Patentin version 3.1 SEQ ID NO 108 LENGTH: 12 TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-990-832C-108
; Sequence 108, Application US/09990832C
; Publication No. US20030149235A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                             , OTHER INFORMATION: Targeting peptide sequence US-09-990-832C-108
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US-10-322-266-434
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Publication No. US20040115135A1
GENERAL INFORMATION:
APPLICANT: Quay, Steven C.
TITLE OF INVENTION: Compositions And Methods For Enhanced Mucosal Delivery Of Peptide
TITLE OF INVENTION: YY And Methods For Treating And Preventing Obesity
FILE REFERENCE: NPC10567
CURRENT APPLICATION NUMBER: US/10/322,266
CURRENT FILING DATE: 2002-12-17
NUMBER OF SEQ ID NOS: 797
SOFTWARE: Patentin version 3.2
LENGTH: 10
TYPE: PRT
               Sequence 4, Application US/10758165
Publication No. US20050196816A1
GENERAL INFORMATION:
APPLICANT: Hammerberg, Bruce
TITLE OF INVENTION: IMMUNOGLOBULIN E DETECTION IN MAMMALIAN SPECIES
FILE REFERENCE: 5051-661
                                                                                                                                                                                                                                                                                 Query Match 35.8%;
Best Local Similarity 62.5%;
Matches 5; Conservative
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CURRENT APPLICATION NUMBER: US/10/758,165
                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
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1; Mismatches
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Pred. No. 4.4e+02
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3.6e+02;
ches 2; Indels
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RESULT 15
US-10-462-452-457
; Sequence 457, Ap
; Publication No.
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Best Local Similarity
6; Conserva
Search completed: May 30,
Job time : 48 secs
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                                                                                                                                                                                  ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-462-452-457
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: PRT; ORGANISM: Ovis aries US-10-758-165-4
                                                                                                                                                                                                                             TITLE OF INVENTION: Compositions and Methods for Enhanced TITLE OF INVENTION: Mucosal-Delivery of Interferon Beta FILE REFERENCE: 02-02US CURRENT APPLICATION NUMBER: US/10/462,452 CURRENT FILING DATE: 2003-06-16 PRIOR APPLICATION NUMBER: 60/393,066 PRIOR RILING DATE: 2002-06-28 NUMBER OF SEQ ID NOS: 790 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 457 LENGTH: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: US 60/440,472
PRIOR FILING DATE: 2003-01-16
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin version 3.2
SEQ ID NO 4
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Best Local
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APPLICANT: El Shafy,
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de Meireles, Jorge
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Pred. No. 5.
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Maximum DB seq
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protein search, using sw model
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length: 15
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1. /EMC Celerra SIDS3/ptodata/1/pubpaa/US09 NEW PUB.pep:*

2. /EMC_Celerra SIDS3/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

3. /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

4. /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US08_NEW_PUB.pep:*

5. /EMC_Celerra_SIDS3/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*

6. /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US10_NEW_PUB.pep:*

7. /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US11_NEW_PUB.pep:*

8. /EMC_Celerra_SIDS3/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
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                     DB
                US-10-538-066-163

US-10-525-043-2

US-11-140-487A-1766

US-10-516-079-74

US-10-516-079-74

US-10-540-431-1

US-11-140-487A-2207

US-11-140-487A-2219

US-11-140-487A-2219

US-11-219-563-93

US-11-219-563-93

US-11-219-563-93

US-11-219-563-93

US-11-140-487A-2219

US-11-140-487A-2219

US-11-1219-563-93

US-11-219-563-93

US-11-219-563-93

US-11-219-335

US-11-176-182-31

US-11-176-182-52

US-11-176-182-53

US-11-176-182-53
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29.481 Million cell updates/sec
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Sequence 2, Appli
Sequence 1766, Ap
Sequence 129, App
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Sequence
Sequence
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                   74, Appl
197, App
2207, Ap
2207, Ap
1810, Ap
1810, Appl
193, Appl
2219, Appl
2219, Appl
2219, Appl
231, Appl
235, Appl
237, Appl
237, Appl
237, Appl
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237, Appl
238, Appl
239, Appl
231, Ap
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US-10-522-043-2
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Best Local S
Matches 4
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| 19 23.5 11 6 US-10-540-431-6 19 23.5 12 1 US-09-784-950-10 19 23.5 12 7 US-11-176-182-112 19 23.5 12 7 US-11-054-072-7056 19 23.5 12 7 US-11-054-072-7059 19 23.5 12 7 US-11-054-072-7069 19 23.5 12 7 US-11-054-072-7060 19 23.5 12 7 US-11-054-072-7061 19 23.5 12 7 US-11-054-072-7061 19 23.5 12 7 US-11-054-072-7063 19 23.5 12 7 US-11-054-072-7063 19 23.5 12 7 US-11-054-072-7063 19 23.5 12 7 US-11-054-072-7065 19 23.5 12 7 US-11-054-072-7066 19 23.5 13 7 US-11-054-072-5136 19 23.5 13 7 US-11-054-072-5136 19 23.5 13 7 US-11-054-072-5138 | 45              | 44           | 43       | 42       | 41                 | 40                 | 39       | 38       | 37       | 36       | 35       | 34       | 33       | 32          | 31       | 30              | 29          | 28           | 27                    |
|---|-----------------|--------------|----------|----------|--------------------|--------------------|----------|----------|----------|----------|----------|----------|----------|-------------|----------|-----------------|-------------|--------------|-----------------------|
| .5 11 6 US-10-540-431<br>.5 12 1 US-07-84-050<br>.5 12 7 US-11-054-072<br>.5 13 7 US-11-054-072<br>.5 13 7 US-11-054-072<br>.5 13 7 US-11-054-072<br>.5 13 7 US-11-054-072   | 19              | 19           | 19       | 19       | 19                 | 19                 | 19       | 19       | 19       | 19       | 19       | 19       | 19       | 19          | 19       | 19              | 19          | 19           | 19                    |
| 6 US-10-540-431 1 US-09-784-950 7 US-11-176-182 7 US-11-054-072   | 23.5            | 23.5         |          |          | •                  |                    |          |          |          |          |          |          |          |             |          |                 |             |              |                       |
| US-11-054-072 US-11-176-182 US-11-176-182 US-11-054-072   | 13              | 13           | 13       | 13       | 12                 | 12                 | 12       | 12       | 12       | 12       | 12       | 12       | 12       | 12          | 12       | 12              | 12          | 12           | 11                    |
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|   | -11-054-072-513 | -11-054-072- | -072-    | 182-     | US-11-054-072-7067 | US-11-054-072-7066 | -11      | -11      |          | -11      | -11      | -11      | -11      | -11-054-072 | -11-054  | -11-054-072-705 | -11-176-182 | -09-784-950- | -10-540- <b>4</b> 31- |
|   | 5138, Ap        | 5137, Ap     | 5136, Ap | 17, App1 | 7067, Ap           | 7066, Ap           | 7065, Ar | 7064, Ap | 7063, Ar | 7062, Ap | 7061, Ar | 7060, Ag | 7059, Ap | 7058, Ap    | 7057, Ag | 7056, Ag        | 112, App    | 10, Appl     | 6, Appli              |

## ALIGNMENTS

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APPLICANT: Epinmune Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7, and TITLE OF INVENTION: Peptides and Compositions FILE REPERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 767
SOFTWARE: PatentIn version 3.2
SEQ ID NO 163
LENGTH: 10
TYPE: PRT
APPLICANT: Lu et al.
TITLE OF INVENTION: Polypeptide
FILE REFERENCE: 69856
CURRENT APPLICATION NUMBER: US/10/522,043
CURRENT FILING DATE: 2005-01-19
PRIOR APPLICATION NUMBER: PCT/GB03/04296
PRIOR FILING DATE: 2003-10-03
PRIOR APPLICATION NUMBER: 0306261.9
PRIOR APPLICATION NUMBER: 0306261.9
PRIOR FILING DATE: 2003-03-19
PRIOR PRIOR PRIOR DATE: 2003-03-19
PRIOR FILING DATE: 2003-03-19
PRIOR FILING DATE: 2003-03-19
PRIOR FILING DATE: 2003-10-07
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                                                                                Sequence 2, Application US/10522043 Publication No. US20060100143A1 GENERAL INFORMATION:
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1; Mismatches
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US-10-525-817-129
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US-11-140-487A-1766
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; OTHER INFORMATION: p53 inhibitor peptide
US-10-522-043-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: hepatitis C virus
US-11-140-487A-1766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/11/140,487A
CURRENT FILING DATE: 2005-05-31
PRIOR APPLICATION NUMBER: EP 04012951.2
PRIOR FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: EP 04447239.7
PRIOR APPLICATION NUMBER: EP 05102441.2
PRIOR PPLICATION NUMBER: EP 05102441.2
PRIOR APPLICATION NUMBER: US 60/576,310
PRIOR APPLICATION NUMBER: US 60/576,310
                                                                       APPLICANT: Nuevolution A/S
TITLE OF INVENTION: Enzymatic encoding
FILE REFERENCE: 9912PC00
CURRENT APPLICATION NUMBER: US/10/525,817
CURRENT FILING DATE: 2005-02-25
NUMBER OF SEQ ID NOS: 165
SOFTWARE: Patentin version 3.1
SEQ ID NO 129
                                                                                                                                                                                                                         Sequence 129, Application US/10525817 Publication No. US20060099592A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/665,395
PRIOR FILING DATE: 2005-03-25
NUMBER OF SEQ ID NOS: 2278
SOFTWARE: Patentin version 3.1
SEQ ID NO 1766
SEQ ID NO 1766
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 28.4%;
Best Local Similarity 57.1%;
Matches 4; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Innogenetics N.V.
TITLE OF INVENTION: Peptides for inducing a CTL and/or HTL response to Hepatitis C
FILE REFERENCE: 166
         LENGTH: 7
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 7
TYPE: PRT
ORGANISM: Artificial Sequence
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   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: EP 05102441.2
FILLING DATE: 2005-03-25
APPLICATION NUMBER: US 60/576,310
FILLING DATE: 2004-06-03
APPLICATION NUMBER: US 60/622,782
FILLING DATE: 2004-10-29
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                                                                                                                                                                                                                                                                                                                                                                 SPLTINO 7
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57.1%;
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; Pred. No. 5.2e
2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 23; DB 6; Length 7; Pred. No. 5.2e+04;
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                                                                                                                                                                                                                                 LENGTH:
10 QYTTTG 15
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FILE REFERENCE: 51530-5006-WO
CURRENT APPLICATION NUMBER: US/10/516,079
CURRENT FILING DATE: 2004-11-29
PRIOR APPLICATION NUMBER: US 60/406,033
PRIOR FILING DATE: 2002-08-27
PRIOR APPLICATION NUMBER: US 60/384,171
PRIOR FILING DATE: 2002-05-31
PRIOR FILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 95
SOFTWARE: PatentIn version 3.2
SEQ ID NO 74
LENGTH: 10
TYPE: PRT
ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                          RESULT 6
US-10-540-431-11
JS-10-540-431-11
Sequence 11, Application US/10540431
Publication No. US20060089307A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Artificially produced US-10-525-817-129
                                                                       US-10-540-431-11
                                                                                                                                                                   FILE REFERENCE: PN0273
; CURRENT APPLICATION NUMBER: US/10/540,431
; CURRENT FILING DATE: 2005-06-22
; NUMBER OF SEQ ID NOS: 14
SOFTWARE: PatentIn version 3.3
SEQ ID NO 11
I.ENCORN: 1
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Best Local Similarity
Matches 4; Conserv
             Query Match
Best Local Similarity
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Matches 4; Conservative
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Publication No. US20060088899A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: ALVAREZ, Vernon L.
APPLICANT: GRIMES, Carol A.
APPLICANT: GONDA, Matchew A.
TITLE OF INVENTION: Combination chemotherapy with chlorotoxin
                                                                                                                                                                                                                                                                                     APPLICANT: Amersham Health AS TITLE OF INVENTION: Contrast /
                                                                                                       ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE: OTHER INFORMATION: Pep21-SCXP_ANDMA sequence
                                                                                      OTHER INFORMATION: Syntetic peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8 TDQYT 12
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Conservative
                27.2%;
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                Score 22; DB 6;
Pred. No. 1.7e+02;
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Pred. No. 5.2e+04;
0; Mismatches 1
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Pred. No. 1.5e+02;
 Mismatches
                                 Length 11;
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/11/140,487A CURRENT FILING DATE: 2005-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            equence 2207, Application US/11140487A
                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: EP 04447239.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 1999-02-08
NUMBER OF SEQ ID NOS: 199
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 09/532,106
PRIOR TILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 09/664,945
PRIOR FILING DATE: 2000-09-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: TRIPED. 7AUC4CP1C
CURRENT APPLICATION NUMBER: US/11/121,282
CURRENT FILING DATE: 2005-05-03
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Innogenetics N.V.
TITLE OF INVENTION: Peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2002-08-30
PRIOR APPLICATION NUMBER: 09/839,666
PRIOR FILING DATE: 2001-04-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 10/372,735
PRIOR FILING DATE: 2003-02-21
PRIOR APPLICATION NUMBER: 10/234,579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Jan-Ingmar Flock
TITLE OF INVENTION: SPECIFICITY EXCHANGERS THAT REDIRECT
TITLE OF INVENTION: ANTIBODIES TO A PATHOGEN
                                                                SOFTWARE: PatentIn version 3.1
                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                        PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: EP 04012951.2
                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 13
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 09/246,258
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APPLICANT: Jan-
Q ID NO 2207
LENGTH: 15
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
OTHER INFORMATION:
                                                                                                                                                                                          FILING DATE: 2005-03-25
APPLICATION NUMBER: US 60/576,310
FILING DATE: 2004-06-03
                                                                                                                              APPLICATION NUMBER: US 60/665,395
                                                                                                                                                                                                                                                        APPLICATION NUMBER: EP 05102441.2
                                                                                                                                                                                                                                                                              FILING DATE: 2004-10-28
                                                                                                                                                  APPLICATION NUMBER: US 60/622,782 FILING DATE: 2004-10-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION:
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b. US20060094862A1
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                                                                                                                                                                                                                                                                                                                                                                                                                            Peptides for inducing a CTL and/or HTL response to Hepatitis C
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80.0%;
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Pred. No. 2e+02;
0; Mismatches
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US-11-140-487A-1810
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US-11-140-487A-2231
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                   PRIOR APPLICATION NUMBER: EP 04012951.2
PRIOR FILING DATE: 2004-06-01
PRIOR PELICATION NUMBER: EP 04447239.7
PRIOR FILING DATE: 2004-10-28
PRIOR FILING DATE: 2004-10-28
PRIOR APPLICATION NUMBER: EP 05102441.2
PRIOR FILING DATE: 2005-03-25
PRIOR APPLICATION NUMBER: US 60/576,310
PRIOR APPLICATION NUMBER: US 60/622,782
PRIOR APPLICATION NUMBER: US 60/622,782
PRIOR APPLICATION NUMBER: US 60/623,782
PRIOR APPLICATION NUMBER: US 60/655,395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-11-140-487A-2231
                                                                                                                                                                                                                                                                                                                                                         Sequence 1810, Application US/11140487A Publication No. US20060093617A1 GENERAL INFORMATION:
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Matches
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Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Publication No. US20060093617A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/11/140,487A CURRENT FILING DATE: 2005-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 60/665,395
PRIOR FILING DATE: 2005-03-25
NUMBER OF SEQ ID NOS: 2278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 166
CURRENT APPLICATION NUMBER: US/11/140,487A
CURRENT FILING DATE: 2005-05-31
                                                                                                                                                                                                                                                                                                                      APPLICANT: Innogenetics N.V.
TITLE OF INVENTION: Peptides for inducing a CTL and/or HTL response to Hepatitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 60/576,310
PRIOR FILING DATE: 2004-06-03
PRIOR APPLICATION NUMBER: US 60/622,782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: EP 04447239.7
PRIOR FILING DATE: 2004-10-28
PRIOR APPLICATION NUMBER: EP 05102441.2
PRIOR FILING DATE: 2005-03-25
PRIOR FILING DATE: 2005-03-25
                                                                                                                                                                                                                                                                                                    FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: EP 04012951.2 PRIOR FILING DATE: 2004-06-01
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TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: hepatitis C virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptides for inducing a CTL
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Pred. No. 2.3e
1; Mismatches
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Pred. No. 2.3e+02;
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1;
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Length 15;

0,

Gaps

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Sequence 19, Application US/10490949

; Publication No. US20060093574A1

; GENERAL INFORMATION:
    APPLICANT: THE REGENTS OF THE UNIVERSITY OF CALIFORNIA

APPLICANT: ALBANI, Salvotore
    APPLICANT: MARTINI, Alberto
    TITLE OF INVENTION: METHODS FOR EPITOPE-SPECIFIC AND CYTOKINE/ANTICYTOKINE
    TITLE OF INVENTION: IMMUNE RESPONSES IN IMMUNE MEDIATED DISEASES
    FILE REFERENCE: UCSD1450-2

; CURRENT APPLICATION NUMBER: US/10/490,949

; CURRENT FILING DATE: 2004-03-25

PRIOR APPLICATION NUMBER: US 60/339,284

PRIOR APPLICATION NUMBER: US 60/339,284

PRIOR APPLICATION NUMBER: US 60/339,284

PRIOR APPLICATION NUMBER: US 60/325,499

PRIOR FILING DATE: 2001-09-25

PRIOR FILING DATE: 2001-09-25

PRIOR FILING DATE: 2001-09-25

NUMBER OF SEQ ID NOS: 59

**CORTENDE: DESCRIPTION OF THE UNIVERSALE AND THE UNIVERSALE AND THE US 60/325,499

**NUMBER OF SEQ ID NOS: 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: MODIFIED ANTIBODIES TO PROSTATE-SPECIFIC TITLE OF INVENTION: MEMBRANE ANTIGEN AND USES THEREOF FILE REFERENCE: 13651.001 (BLI-001)
CURRENT APPLICATION NUMBER: US/11/219,563
CURRENT FILING DATE: 2005-09-02
PRIOR APPLICATION NUMBER: PCT/US04/06586
PRIOR FILING DATE: 2004-03-03
PRIOR APPLICATION NUMBER: US/1379,838
PRIOR APPLICATION NUMBER: US/379,838
PRIOR FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: US/379,838
PRIOR FILING DATE: 2003-05-30
PRIOR FILING DATE: 2003-05-30
NUMBER: PSTATION NUMBER: 10/449,379
PRIOR FILING DATE: 2003-05-30
NUMBER: DATE: 2003-05-30
NUMBER: DATE: 2003-05-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; LENGTH: 10
; TYPE: PRT
; ORGANISM: Mus musculus
US-11-219-563-93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                밁
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; TYPE: PRT
; ORGANISM: hepatitis C virus
US-11-140-487A-1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-490-949-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 93, Applica Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 2278
SOFTWARE: PatentIn version
SEQ ID NO 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bander,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11 YTTTG 15
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PatentIn version 3.1
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Pred. No. 5.2e+04;
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2.3e+02;
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US-10-489-071-6
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US-11-140-487A-2219
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/11/140,487A
CURRENT FILING DATE: 2005-05-31
PRIOR APPLICATION NUMBER: EP 04012951.2
PRIOR FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: EP 04447239.7
PRIOR FILING DATE: 2004-10-28
PRIOR PPLICATION NUMBER: EP 05102441.2
PRIOR APPLICATION NUMBER: EP 0505441.2
PRIOR APPLICATION NUMBER: US 60/576,310
PRIOR APPLICATION NUMBER: US 60/576,310
PRIOR FILING DATE: 2004-06-03
PRIOR FILING DATE: 2004-06-03
                                  Sequence 6, Application US/10489071

Publication No. US20060094672A1

GENERAL INFORMATION:
APPLICANT: PASQUALINE ET AL.

TITLE OF INVENTION: COMPOSITIONS AND METHODS OF USE OF TARGETING PEPTIDES AGAINST

TITLE OF INVENTION: PLACENTA AND ADIPOSE TISSUES

FILE REFERENCE: UTSC: 856US

CURRENT APPLICATION NUMBER: US/10/489,071

CURRENT FILING DATE: 2004-03-08

PRIOR APPLICATION NUMBER: PCT/US02/27836

PRIOR FILING DATE: 2002-08-30

NUMBER OF SEQ ID NOS: 144

SEQ ID NO 6

LENGTH: 7
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LENGTH: 15
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SEQ ID NO 2219
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Best Local Similarity 57.18
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/622,782
PRIOR FILING DATE: 2004-10-29
PRIOR APPLICATION NUMBER: US 60/665,395
PRIOR FILING DATE: 2005-03-25
NUMBER OF SEQ ID NOS: 2278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Innogenetics N.V.
TITLE OF INVENTION: Peptides for inducing a
FILE REFERENCE: 166
                   TYPE: PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
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80.0%;
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Pred. No. 3.5e+02;
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Pred. No. 3.5e+02
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US-10-525-817-130

US-10-525-817-130

Sequence 130, Application US/10525817

Publication No. US20060099592A1

GENERAL INFORMATION:
APPLICANT: Nuevolution A/S

TITLE OF INVENTION: Enzymatic encoding
FILE REFERENCE: P912PC00

CURRENT APPLICATION NUMBER: US/10/525,817

CURRENT FILING DATE: 2005-02-25

NUMBER OF SEQ ID NOS: 165

SOFTWARE: PatentIn version 3.1

SEQ ID NO 130

LENGTH: 7

TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Artificially produced

US-10-525-817-130
Search completed: May 30, 2006, 10:25:04 Job time: 5.66667 secs
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; OTHER INFORMATION: Synthetic Peptide
US-10-489-071-6
                                                                                                                                                     Query Match 24.7%; Score 20; DB 6; Length 7; Best Local Similarity 80.0%; Pred. No. 5.2e+04; Matches 4; Conservative 0; Mismatches 1; Indels
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Result
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Maximum Match 100%
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    length:
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Match
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1: /EMC Celerra SIDS3/ptodata/2/iaa/5_COMB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/2/iaa/6_COMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/iaa/7_COMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/iaa/H_COMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/iaa/PCTUS_COMB.pep:*

6: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*

7: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*
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Gapop 10.0 , Gapext 0.5
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                                        GenCore version 5.1.8 (c) 1993 - 2006 Biocceleration Ltd
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US-09-641-528B-35442
US-09-641-528B-507
US-09-641-528B-51014
US-09-641-528B-51017
US-09-641-528B-10417
US-09-641-528B-10417
US-09-641-528B-10417
US-09-641-528B-10362
US-09-641-528B-10362
US-09-641-528B-10362
US-09-641-528B-47375
US-09-641-528B-47375
US-09-641-528B-47375
US-09-641-528B-47375
US-09-641-528B-50683
US-09-341-982-78
US-09-341-982-78
US-09-341-982-78
US-09-341-528B-50683
US-09-341-528B-50683
US-09-341-529-161
              US-09-641-528B-4318
US-09-641-528B-50956
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                                                                                                                         Sequence 29, Appl
Sequence 48206, A
Sequence 5634, Ap
Sequence 10412, A
Sequence 32442, A
Sequence 51014, A
Sequence 51014, A
Sequence 5507, Ap
Sequence 5507, Ap
Sequence 35452, A
Sequence 35452, A
Sequence 10362, A
Sequence 10363, A
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50683, Appl
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161, App
1418, Ap
50956, A
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29, Appl
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| 45                   | 44                  | 43              | 42              | 41               | 40              | 39              | 38                  | 37                   | 36                   | 35                   | 34               | သ                    | 32              | 31                | 30               | 29              | 28              | 27                   |
|----------------------|---------------------|-----------------|-----------------|------------------|-----------------|-----------------|---------------------|----------------------|----------------------|----------------------|------------------|----------------------|-----------------|-------------------|------------------|-----------------|-----------------|----------------------|
| 25                   | 25                  | 25              | 25              | 25               | 25              | 25              | 25                  | 25                   | 25                   | 25                   | 25               | 26                   | 26              | 26                | 26               | 26              | 26              | 26                   |
| 30.9                 | 30.9                | 30.9            | 30.9            | 30.9             | 30.9            | 30.9            | 30.9                | 30.9                 | 30.9                 | 30.9                 | 30.9             | 32.1                 | 32.1            | 32.1              |                  | 32.1            | 32.1            | 32.1                 |
| 11                   | 11                  | 11              | 11              | 11               | 11              | 11              | 9                   | 80                   | œ                    | œ                    | 00               | 15                   | 15              | 12                | 12               | 12              | 12              | 11                   |
| ω                    | W                   | Н               | ۲               | ₽                | μ.              | 4               | w                   | ω                    | w                    | w                    | ۳                | w                    | N               | ω                 | N                | N               | Н               | ω                    |
| US-09-641-528B-14203 | US-09-641-528B-5591 | US-08-465-334-2 | US-08-461-621-2 | US-07-949-812-24 | US-08-461-599-2 | US-08-077-939-2 | US-09-641-528B-5590 | US-09-641-528B-42396 | US-09-641-528B-35388 | US-09-641-528B-10393 | US-08-413-708B-4 | US-09-641-528B-50625 | US-10-378-707-1 | US-10-279-991A-13 | US-09-517-866-13 | US-09-514-739-5 | US-08-423-441-4 | US-09-641-\$28B-4319 |
| Sequence 1           | Sequence 51         | Sequence 2      | Sequence 2      | Sequence 24,     | Sequence 2      | Sequence 2      | Sequence 59         | Sequence 4:          | Sequence 3:          | Sequence 1           | Sequence 4       | Sequence 5           | Sequence 1      | Sequence 13       | Sequence 1:      | Sequence 5      | Sequence 4      | Sequence 4:          |
| 14203, A             | 5591, Ap            | , Appli         | , Appli         | 4, Appl          | , Appli         | , Appli         | 5590, Ap            | 42396, A             | 35388, A             | 10393, A             | 4, Appli         | 50625, A             | , Appli         |                   | 3, Appl          | Appli           | Appli           | 4319, Ap             |

ALIGNMENTS

#### GENERAL INFORMATION: APPLICANT: FOX, GARY M. APPLICANT: JING, SHUGIAN APPLICANT: WEN, DUANZHI ITITLE OF INVENTION: GLIAL CELL LINE-DERIVED NEUROTROPHIC FACTOR RECEPTOR FILE REFERENCE: A-401C CURRENT APPLICATION NUMBER: US/08/837,199A CURRENT FILING DATE: 1997-044-14 PRIOR APPLICATION NUMBER: US 60/015,907 PRIOR APPLICATION NUMBER: US 60/015,907 PRIOR FILING DATE: 1996-04-22 PRIOR FILING DATE: 1996-04-22 PRIOR FILING DATE: 1996-05-09 NUMBER OF SEQ ID NOS: 47 SOFTWARE: Patentin version 3.1 SEQ ID NO 29 LENGTH: 14 TYPE: PRT RESULT 1 5223254-6 밁 Ş ; ORGANISM: RAT US-08-837-199A-29 US-08-837-199A-29 5223254-6 PATENT NO. 5223254 APPLICANT: PARADISO, PETER R.; HILDRETH, STEPHEN W.; HU, BRANDA T.; MARTIN-GALLARDO, ANTONIA; ARUMUGHAM, RASAPPA TITLE OF INVENTION: RESPIRATOTY SYNCYTIAL VIRUS VACCII NUMBER OF SEQUENCES: 17 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/07/247,017 FILING DATE: 20-SEP-1988 PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION UMBER: 102,180 SEQ ID NO:6: Sequence 29, Application Patent No. 6455277 Matches Query Match FILING DATE: 29-SEP-1987 Local LENGTH: 13 N 2 NDSPIQTDQ 10 Similarity 6; Conserv NDMPITNDO 10 Conservative 39.5**%**; 66.7**%**; US/08837199A Score 32; DB Pred. No. 36; Mismatches SYNCYTIAL VIRUS VACCINES DB 7; Length 13; Indels 0 Gaps

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RESULT 4
US-09-641-528B-5634
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                                                                   APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esteban
APPLICANT: Crey, Howard
TITLE OF INVENTION: USING PAPTIDE AND NUCLEIC ACID COMPOSITIONS
TITLE OF INVENTION: USING PAPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5634
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS TITLE OF INVENTION: USING PEFTIDE AND NUCLEIC ACID COMPOSITIONS FILE REFERENCE: 2060.0100001

CURRENT APPLICATION NUMBER: US/09/641,528B

CURRENT FILING DATE: 2000-08-15

PRIOR APPLICATION NUMBER: US 60/172,705

PRIOR PEPILONION NUMBER: US 60/172,705

PRIOR FILING DATE: 1999-12-10

NUMBER OF SEQ ID NOS: 51505

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 48206

LENGTH: 15
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Best Local Similarity 60.
Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 5634, Application US/09641528B Patent No. 7026443
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APPLICANT:
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                                                           LENGTH: 9
OTHER INFORMATION: Peptide Derived from Human Papillomavirus
                     ORGANISM: Artificial Sequence FEATURE:
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Celis, Esteban
Grey, Howard
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Pred. No. 1.
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Pred. No. 58;
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Sequence 35442, Application US/09641528B

| Sequence 35442, Application US/09641528B
| Patent NO. 7026443|
| GENERAL INFORMATION:
| APPLICANT: Sette, Alessandro
| APPLICANT: Sette, Alessandro
| APPLICANT: Southwood, Scott
| APPLICANT: Chesnut, Robert
| APPLICANT: Chesnut, Robert
| APPLICANT: Celis, Esteban
| APPLICANT: Grey, Howard
| TITLE OF INVENTION: INDUCING CELLULAR INMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
| TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
| FILE REFERENCE: 2060.0100001
| CURRENT APPLICATION NUMBER: US/09/641,528B
| CURRENT APPLICATION NUMBER: US/09/641,528B
| CURRENT FILING DATE: 2000-08-15
| PRIOR APPLICATION NUMBER: US 60/172,705
| PRIOR FILING DATE: 1999-11-10
| NUMBER OF SEQ ID NOS: 51505
| SOFTMARE: FastSEQ for Windows Version 4.0
| LENGIN ON 35442
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US-09-641-528B-35442
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APPLICANT: Carey, Howard
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PETTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
SPRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 10412
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GENERAL INFORMATION:
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Best Local
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                  LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Peptide Derived from Human Papillomavirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence FEATURE:
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FEATURE:
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Local Similarity 50.0%;
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Chesnut, acc
Chesnut, acc
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Pred. No. 5e+(
2; Mismatches
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Pred. No. 5e+05;
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5e+05;
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US-09-641-528B-51014; Sequence 51014, Application US/09641528B
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US-09-641-528B-42435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 42435
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                                           APPLICANT: Celis. Beteban
APPLICANT: Cely, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 51014
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APPLICANT:
APPLICANT:
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APPLICANT:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                  ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                    INFORMATION:
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Grey, Howard
                                                                                                                                                                                                                                                             Chesnut, Robert
                                                                                                                                                                                                                                                                             Sidney, John Southwood, So
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Southwood,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chesnut,
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Pred. No. 5e+05;
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; FEATURE: ; OTHER INFORMATION: Peptide Derived from Human Papillomavirus US-09-641-528B-51014
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US-09-641-528B-10417
; Sequence 10417, Application US/09641528B
; Patent No. 7026443
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TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10417
LENGTH: 10
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5507
LENGTH: 10
TYPE: PRT
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Best Local Similarity
Matches 4; Conserv
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PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
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APPLICANT: Grey, Howat
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,5288
CURRENT FILING DATE: 2000-08-15
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5. 7026443
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Similarity 50.0%;
4; Conservation
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50.0%;
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Pred. No. 1.3e+02;
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APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esteban
APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FASTSEQ for Windows Version 4.0
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TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR PILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 35452
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APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
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Best Local Similarity 50.0
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APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
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Pred. No.
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Pred. No. 1.3e+02
2; Mismatches
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US-09-641-528B-10362
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US-09-641-528B-5635
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APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esceban
APPLICANT: Grey, Howard
TITLE OF INVENTION: UNDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.010001
CURRENT FILING DATE: 2000-08-15
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10362
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PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 5635
                                                                                                                                                                                                                                                                                                                                           Patent No. 7026443
GENERAL INFORMATION:
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Best Local Similarity
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Patent No. 702644
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APPLICANT: Sidney, John
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.0100001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT APPLICATION ALTREES 106.017.765
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APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
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ORGANISM: Artificial Sequence
FEATURE:
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Pred. No. 1.5e+02;
2; Mismatches 2
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Pred. No. 1.3e+02
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APPLICANT: Sidney, John
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Cols, Esteban
APPLICANT: Celis, Esteban
APPLICANT: Celis, Esteban
APPLICANT: Grey, Howard
TITLE OF INVENTION: INDUCING CELLULAR IMMUNE RESPONSES TO HUMAN PAPILLOMAVIRUS
TITLE OF INVENTION: USING PEPTIDE AND NUCLEIC ACID COMPOSITIONS
FILE REFERENCE: 2060.010001
CURRENT APPLICATION NUMBER: US/09/641,528B
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR APPLICATION NUMBER: US 60/172,705
PRIOR FILING DATE: 1999-12-10
NUMBER OF SEQ ID NOS: 51505
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 35263
LENGTH: 11
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Peptide Derived from Human Papillomavirus
US-09-641-528B-35263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; LENGTH: 11
; TYPE: PRT
; ORGANISM: Artificial Sequence
; EBATURE:
; FEATURE:
; OTHER INFORMATION: Peptide Derived from Human Papillomavirus
US-09-641-520B-10362
Search completed: May 30, 2006, 11:11:07 
Job time : 21.333 secs
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                                                                                                                                                                        Query Match
Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 34.6%;
Best Local Similarity 50.0%;
Matches 4; Conservative
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APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Chesnut, Robert
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                                                                                                                                                                        Score 28; DB 3; Length 11; 
Pred. No. 1.5e+02; 
2; Mismatches 2; Indels
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Pred. No. 1.5e+02;
2; Mismatches 2; Indels
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Maximum DB seq
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 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Match
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123.707 Million cell updates/sec
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Listing first 45
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Copyright (c) 1993 - 2006 Biocceleration Ltd
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| 45                 | 44                | 43                 | 42                 | 41                 | 40                 | 39                 | 38                 | 37                 | 36                 | 35                | 34                 | ω<br>ω             | 32                 | 31                 | 30                 |
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| 21.0               | 21.0              | 21.0               | 21.0               | 21.0               | 21.0               | 21.0               | 21.0               | 21.0               | 21.0               | 21.0              | 22.2               | 22.2               | 22.2               | 22.2               | 22.2               |
| 12                 | 12                | 12                 | 12                 | 11                 | 11                 | 11                 | 10                 | 9                  | 9                  | ထ                 | 15                 | 15                 | 15                 | 15                 | 14                 |
| N                  | N                 | N                  | N                  | N                  | N                  | N                  | N                  | N                  | N                  | N                 | 2                  | N                  | N                  | N                  | N                  |
| PN0170             | JU0356            | PT0228             | S25485             | \$42449            | A29806             | C53652             | E61512             | PL0139             | PT0247             | 157018            | S72432             | S29485             | PA0097             | S32677             | PT0232             |
| alcohol dehydrogen | cycloleonurinin - | Ig heavy chain CDR | transcription fact | ant1 protein - pha | acidic proline-ric | rhlR protein - Pse | variant surface gl | carbon-monoxide de | Ig heavy chain CRD | gene Cftr protein | epoxypropan isomer | GTP-binding protei | starch phosphoryla | nitrogenase cofact | Ig heavy chain CRD |

# ALIGNMENTS

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Ig heavy chain DJ region (clone C96-100) - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1306
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
U, Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymph
A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1306
                                                                                                                                                                                                                                               R;Frebort, I.; Tamaki, H.; Ishida, H.; Pec, P.; Luhova, L.; Tsuno, H.; Halata, M.; Asano Eur. J. Biochem. 237, 255-265, 1996
A;Title: Two distinct quinoprotein amine oxidases are induced by n-butylamine in the myc A;Reference number: S71303; MUID:96203933; PMID:8620882
A;Accession: S71304
                                                                                                                                                                                                                                                                                                                                                        amine oxidase (copper-containing) (EC 1.4.3.6) II - Aspergillus niger (fragment) C;Species: Aspergillus niger (C;Date: 12-Feb-1998 #sequence_revision 01-May-1998 #text_change 09-Jul-2004 C;Accession: S71104
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S71304
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C;Keywords: copper binding;
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A;Residues: 1-14 <WAS>
A;Crose-references: UNIPARC:UPI000017C253
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                           A; Molecule type: protein A; Residues: 1-11 <FRE>
                                                                                  Matches
                                                                                                         Query Match
Best Local
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5; Conserv
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NDSPALND
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                                                                                    Conservative
                                                                                                                                                                   UNIPROT:Q7M504; UNIPARC:UPI000017B3B7
binding; monomer; oxidoreductase; quinoprotein;
                                                                                                    32.1%;
62.5%;
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                                                                              Score 26; DB 2; Lei
Pred. No. 1.6e+02;
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RESULT 3 PS0249

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R;Tsugita, A.; Kamo, M.
submitted to JIPID, April 1993
A;Reference number: PS0209
A;Accession: PS0259
A;Molecule type: protein
A;Residues: 1-11 <TSU>
                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 5
PS0259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ig heavy chain DJ region (clone C85-1B) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1305
C;Accession: PH1305
R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
J. Exp. Med. 176, 1577-1581, 1992
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A;Accession: PH1305
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submitted to JIPID, April 1993
A;Reference number: PS0206
A;Accession: PS0249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           porin - rice (strain Nihonbare) (fragment)
C;Species: Oryza sativa (rice)
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 09-Jul-2004
C;Accession: PS0249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: DNA
A; Residues: 1-14 <WAS>
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A;Experimental source: callus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: protein
A; Residues: 1-14 <TSU>
                                                                                                                                                                                                                                                                                                                ;Species: Oryza sativa (rice);Date: 03-Feb-1904 #sequence_revision 03-Feb-1904 #...;Pate: 03-Feb-1904 #...;Pate: PS0259
                                                                                                                                                                                        ;Cross-references: UNIPARC:UPI000017B102;Experimental source: callus
                                                                                                                      Query Match
Best Local
                                                                                                                                                                      Comment: molecular weight 39K, pI 5.7.
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Best Local :
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Best Local
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Similarity 50.0%;
4; Conservative
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57.1%;
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Pred. No.
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1.1e+03;
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1.3e+03
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RESULT 6

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RESULT 8
S04875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            urinary tract stone matrix protein 10, 42K - human (fragment) C;Species: Homo sapiens (man) C;Date: 12-Apr-1995 #sequence_revision 12-Apr-1995 #text_change C;Accession: H56046 R;Binette, J.P.; Binette, M.B.; Gawinowicz, M.A.; Kendrick, N. submitted to the Protein Sequence Database, February 1995
                                                                                                                                                                                                                          R;Ebeling, S. submitted to the EMBL Data A;Reference number: S04873 A;Accession: S04875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         T-cell antigen receptor VJ junction beta chain - C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-199
C;Accession: S47376
R;Lehner, P.J.
                                                                                                                                                                                                                                                                                                nifs protein - Bradyrhizobium japonicum (fragment) C;Species: Bradyrhizobium japonicum C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 C;Accession: S04875
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                                                                                                                        A; Gene: nifS
A; Start codon: GTG
                                                                                                                                                           A;Cross-references: UNIPROT:P37030; UNIPARC:UPI000016E719; EMBL:X13691; NID:g39544; C;Genetics:
                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-11 <EBE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: UNIPARC:UPI0000116687; EMBL:Z35702; NID:g527497; C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-13 <LEH>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     submitted to the EMBL Data Library, August 1994
A;Description: Human HLA-A0201 restricted recognition
A;Reference number: $47355
A;Accession: $47376
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A; Accession: H56046
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2 NDSPIQTD
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4; Conserv
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50.0%;
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                                                                     Score 20; DB 2;
Pred. No. 1.9e+03;
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Pred. No. 1.9e+03;
0; Mismatches 2
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Pred. No. 1.5e+03;
1; Mismatches 3
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                                                      Mismatches
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T-cell receptor beta chain (PES.1.1) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 05-Nov-1999
C;Accession: PHO771
R;Casanova, J.L.; Romero, P.; Widmann, C.; Kourilsky, P.; Maryanski, J.L.
J. Exp. Med. 174, 1371-1383, 1991
A;Title: T cell receptor genes in a series of class I major histocompatibility complex-1
allelic exclusion and antigen-specific repertoire.
A;Reference number: PHO746; MUID:92078846; PMID:1836010
A;Accession: PHO771
A;Molecule type: mRNA
A;Residues: 1-12 <CAS-
A;Reference mRNA
A;Residues: 1-12 <CAS-
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S05002
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A;Title: Ascaris suum phosphofructokinase. Phosphorylation by protein kinase and A;Reference number: A26120; MUID:87083467; PMID:3025208
A;Accession: A26120
                                                                                                                                                                                                                                                                                                                                RESULT 11
PH0771
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C;Keywords: amidated carboxyl end; neuropeptide; pyroglutamic acid
F;11/Modified site: pyrrolidone carboxylic acid (Cln) #status experimental
F;11/Modified site: amidated carboxyl end (Asn) #status experimental
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C;Date: 07-Sep-1990 #sequence_revision 09-Apr-1998 #text_change 09-Jul-2004
C;Accession: S05002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: protein A; Residues: 1-11 < KUL>
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Cross-references: UNIPARC:UPI0000115FBE; EMBL:X60865; NID:g53624; PIDN:CAA43255.1; PID
Experimental source: T lymphocyte
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Similarity 50.0%;
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Pred. No.
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Pred. No. 1.9e+03;
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1.9e+03;
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C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1347
C;Accession: R; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
J. Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymph A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1347
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PH1347
                                                                                               leucosulfakinin-II - Madeira cockroach
N;Alternate names: LSK-II
C;Species: Leucophaea maderae (Madeira cockroach)
C;Date: 31-Dec-1988 #sequence_revision 31-Dec-1988 #text_change 09-Jul-2004
C;Accession: A26335
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A;Title: Glu, Asp-specific proteinase from Actinomycetes.
A;Reference number: PN0125
A;Accession: PN0125
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PN0125
                 R;Nachman, R.J.; Holman, G.M.; Cook, B.J.; Haddon, W.F.; Ling, N. Biochem. Biophys. Res. Commun. 140, 357-364, 1986
A;Title: Leucosulfakinin-II, a blocked sulfated insect neuropeptide with homology A;Reference number: A26335; MUID:87048769; PMID:3778455
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A;Note: article in Russian with English abstract
C;Keywords: hydrolase; serine proteinase
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A; Residues: 1-13 < MOS>
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C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 09-Jul-2004
A; Reference number: A; Accession: A26335
                                                                                                                                                                                                                             RESULT 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPARC:UPI000017C21CC;Keywords: heterotetramer; immunoglobulii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-14 <WAS>
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A; Molecule type: protein
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A; Cross-references: UNIPROT: P09039; UNIPARC: UPI000012E961
A; Cross-references: UNIPROT: P09039; UNIPARC: UPI000012E961
C; Comment: This peptide was isolated from head extracts. It stimulates muscle contractic C; Superfamily: gastrin
C; Keywords: amidated carboxyl end; neuropeptide; pyroglutamic acid; sulfoprotein
F; 1/Modified site: pyrrolidone carboxylic acid (cln) #status experimental
F; 5/Binding site: sulfate (Tyr) (covalent) #status experimental
F; 10/Modified site: amidated carboxyl end (Phe) #status experimental
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C;Keywords: amidated carboxyl end; neuropeptide; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;10/Modified site: amidated carboxyl end (Phe) #status experimental
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A;Title: Isolation and structure of two gastrin/CCK-like neuropeptides from the American A;Reference number: A60656; MUID:90137190; PMID:2615921
A;Accession: B60556
A;Molecule type: protein
A;Residues: 1-10 <VEE>
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STRAIN=IC10, IC2, IC3, IC4, IC5, IC6, IC7, IC8, and I PubMed=15956661; DOI=10.1534/genetics.105.045120; Borge T., Webster M.T., Andersson G., Saetre G.-P.; "Contrasting Patterns of Polymorphism and Divergence Chromosome and Autosomes in Two Ficedula Flycatcher S
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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PubMed=15956661; DOI=10.1534/genetics.105.045120
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IL; AJ869946; CAI35699.1; -; Genomic_DNA.
IL; AJ869947; CAI35700.1; -; Genomic_DNA.
IL; AJ869948; CAI35701.1; -; Genomic_DNA.
IL; AJ869949; CAI35702.1; -; Genomic_DNA.
IL; AJ869950; CAI35703.1; -; Genomic_DNA.
IL; AJ869951; CAI35704.1; -; Genomic_DNA.
IL; AJ869952; CAI35705.1; -; Genomic_DNA.
IL; AJ869952; CAI35705.1; -; Genomic_DNA.
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1L; AJ869957; CAI35710.1; -; Genomic_DNA.
1L; AJ869959; CAI35711.1; -; Genomic_DNA.
1L; AJ869959; CAI35712.1; -; Genomic_DNA.
1L; AJ869960; CAI35713.1; -; Genomic_DNA.
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Ficedula parva (Red-breasted flycatcher).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Passeriformes; Muscicapidae; Ficedula.
                                                          Tsugita A.;
Submitted (APR-1993) to
                                                                                                                                                                                 Eukaryota; Viridiplantae; Stre
Spermatophyta; Magnoliophyta;
Ehrhartoideae; Oryzeae; Oryza.
NCBI TaxID=4530;
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Primmer C.R., Syvanen A
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TISSUB-Rectal gland;
TISSUB-Rectal gland;
MEDLINE=21534268; PubMed=11676495; DOI=10.1006/bbrc.2001.5826;
Schuurmans Stekhoven F.M.A.H., Flik G., Wendelaar Bonga S.E.;
"N-rerminal sequences of small ion channels in rectal glands o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Viridiplantae; Streptophyta;
Spermatophyta; Magnoliophyta; eudicotylo
rosids; eurosids I; Fabales; Fabaceae; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=91378472; PubMed=1898034; Chilson O.P., Kelly-Chilson A.E., S "Pyrroline-5-carboxylate reductase
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Q9S929;
                                                                                                                                                                                                                                                                                                 Lamna nasus (Porbeagle) (Squalus nasus).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Galeomorphii; Galeoidea; Lamniformes; Alopiidae;
                                                                                                                                                                                                                                                                                                                                                                                                    29-MAR-2005, integrated into Unipol-
01-OCT-2001, sequence version 1.
07-FEB-2006, entry version 16.
Phospholemman-like protein (PLMS)
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MEDLINE=91378472;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STANDARD;
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Pred. No. 5e+0
1; Mismatches
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Pred. No. 5.4e+03
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Cell wall
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Differential extraction and protein sequencing differences in patterns of primary cell wall pro J. Biol. Chem. 272:15841-15848 (1997).
-i- SUBCELLULAR LOCATION: Cell wall.
-i- SIMILARITY: Belongs to the thaumatin family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROTEIN SEQUENCE, AND SUBCELLULAR LOCATION.
MEDLINE=97332671; PubMed=9188482; DOI=10.1074/jbc.272.25.15841;
Robertson D., Mitchell G.P., Gilroy J.S., Gerrish C., Bolwell G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P80781;
11-OCT-2005, integrated into Unil
11-OCT-2005, sequence version 1.
07-MAR-2006, entry version 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nicotiana tabacum (Common tobacco).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnaliophyta; eudicotyledons; core eudicotyledons;
asterids; lamiids; Solanales; Solanaceae; Nicotiana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
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-!- FUNCTION: Induces a hyperpolarization-activated chloride current when expressed in Xenopus oocytes. May have a functional role in
                                                                                                                                                                                                         CHAIN
                                                                                                                                                                                                                                                                                                                                                                        Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                                              Copyrighted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=4097,
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                                                                                                                                                                                                                                                                                                     InterPro; IPR001938; Thaumatin
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SUBCELLULAR LOCATION: Type I membrane protein.
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SIMILARITY: Belongs to the FXYD family.
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>15 Osmotin-like
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1744 MW;
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O_0000096233.
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MEDLINE=98223151; PubMed=9556557; DOI=10.1074/jbc.273.18.10843;
Di Noia J.M., D'Orso I., Aslund L., Sanchez D.O., Frasch A.C.;
"The Trypanosoma cruzi mucin family is transcribed from hundred genes having hypervariable regions.";
genes having hypervariable regions.";
J. Biol. Chem. 273:10843-10850(1998).
                                 "Proteomics of the venom from the Amazonian scorpion Tityus and the role of prolines on mass spectrometry analysis of to J. Chromatogr. B 803:55-66(2004).

-i- SUBCELLULAR LOCATION: Secreted protein.
-i- TISSUE SPECIFICITY: Expressed by the venom gland.
-i- MASS SPECTROMETRY: MW=7109.4; METHOD=Electrospray; RANGE
                                                                                                                                                                                                           Tityus cambridgei (Amazonian scorpion).
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Scorpiones;
Buthida; Buthoidea; Buthidae; Tityus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mucin-like protein
Name=EMUCe-19c8;
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Q9TVF1;
Copyrighted by
                                                                                                           PubMed=15025998; DOI=10.1016/j.jchromb.2003.09.002;
Batista C.V.F., del Pozo L., Zamudio F.Z., Contrera
Wanke E., Possani L.D.;
                                                                                                                                                              SPECTROMETRY
                                                                                                                                                                        PROTEIN SEQUENCE,
                                                                                                                                                                                              NCBI_TaxID=184226;
                                                                                                                                                                                                                                                 Toxin Tc41 (Fragment)
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NCBI_TaxID=5693;
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                                                                                                                                                                          SUBCELLULAR LOCATION, TISSUE SPECIFICITY,
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80.0%;
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Pred. No. 2.8e+06;
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RESULT 12
Q54394 STRLI
Q54394;
ID Q54394;
AC Q54394;
DT 01-NOV-1996, integrated into UniProtKB/
DT 01-NOV-1996, sequence version 1.
DT 07-FEB-2006, entry version 19.
DE S-adenosyl-L-homocysteine hydrolase (From Streptomyces lividans, actinobacteria; Actinobacteria; Actinobacteria; RT (1)
RN (1)
RP NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STAN COCCEPTION AND COCCEPTION OF THE COCCEPTION
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Q71GS6_9HYME
ID Q71GS6_9HYME
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05-JUL-2004,
07-MAR-2006,
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PubMed=16343953; DOI=10.1016/j.ympev.2005.10.003; Larkin L.L., Neff J.L., Simpson B.B.; Phylogeny of the Callandrena subgenus of Andrena Andrenidae) based on mitochondrial and nuclear DNI convergent evolution.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-JUL-2004, sequence version 1.
07-MAR-2006, entry version 8.
Cytochrome oxidase subunit I (Fragment).
                                                                            Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales; Streptomycineae; Streptomycetaceae; Streptomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; AF504376; AAQ07723.1; -; Genomic_DNA.GO; GO:0005739; C:mitochondrion; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Arthropoda;
Neoptera; Endopterygota; Hymenop
Andrenidae; Andreninae; Andrena;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
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NON_TER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mol. Phylogenet. Evol. 38:330-343(2006).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=205171;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DOI=10.1016/j.ympev.2005.10.003; f J.L., Simpson B.B.;
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; Hymenoptera; Apocrita; Aculeata; Apo
Andrena; Callandrena.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 22; DB 2;
Pred. No. 1.1e+04;
1; Mismatches 2
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Pred. No. 7.8e+03;
}; Mismatches 2;
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/FTId=PRO_0000066806.
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data: Polyphyly
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RESULT 13
UC19 MAIZE
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DT 01-F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 genome analysis program.";
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01-OCT-1996, integrated into Unit
01-OCT-1996, sequence version 1.
07-FEB-2006, entry version 21.
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                                                                                                                                                                                                                                                                                                                                                                                                              Distributed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Theor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Touzet P., Riccardi F., Morin C., Damerval C., Huet J.-C., Pernollet J.-C., Zivy M., de Vienne D.; "The maize two dimensional gel protein database: towards an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; X79814; CAA56211.1; -; Genomic_DNA. GO; GO:0004013; F:adenosylhomocysteinase activity; IEA. GO; GO:0016787; F:hydrolase activity; IEA. GO; GO:0016730; P:one-carbon compound metabolism; IEA.
                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Theor. Appl. Genet. 93:997-1005(1996).
-!- MISCELLANEOUS: On the 2D-gel the determined protein is: 5.6, its MW is: 18.4 kDa.
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Pfam; PF05221; AdoHcyase; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEMS Microbiol. Lett. 1217:79-84(1995).
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"Activation of the actinorhodin biosynthetic pathway in Streptomyces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=TK21;
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3; Conservative
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                                          QYTTTG 15
  RYTTYG
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7 AA;
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                                                                                                                                                                                                                                                                                              sequencing.
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15
1672 MW;
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33.3%;
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2D-PAGE of etiolated coleoptile (Spot 406)
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; Pred. No. 1.1e.
2; Mismatches
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                                                                                     Score 22; DB 1;
Pred. No. 1.2e+04;
l; Mismatches 1
                                                                                                                                                                                                                                         Unknown protein from 2D-PAGE coleoptile. /FTId=PRO_0000055515.
                                                                                                                                                                              1CF69D4DA8737F9D CRC64;
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1.1e+04;
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                                                                                                                                Length 15;
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ID Q71138;
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DE Cytochrome oxi
OS Andrena auripe
OG Mitochondrion.
OC Eukaryota; Mett
OC Neoptera; Endo
OC Andrenidae; An
OX NCBI_TaxID=205
RN [1]
RP NUCLEOTIDE SEQ
RC STRAIN=126;
RX PUBMEd-1634395
RA Larkin L.L.; N
RT "Phylogeny of
RT Andrenidae) ba
RT convergent evo
RL MG1. Phylogene
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Neoptera; Endopterygota; Hymenoptera; Apocrit
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   EMBL; AF504320;
GO; GO:0005739;
                                            Copyrighted by the UniProt Consortium, see http://www.uniprot.org/t.Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
STRAIN=126;
                                                                                                                                                                                                                                                                                                     Mitochondrion.
                                                                                                                                                                                                                                                                                                                  Andrena auripes.
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Cytochrome oxidase subunit I (Fragment).
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C:mitochondrion;
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ptera; Apocrita; Acu
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1 data: Polyphyly
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